

Result No.	Query			DB	ID	Description
	Score	Match	Length			
1	1431	100.0	282	4	US-09-404-879A-393	Sequence 393, Appl
2	1431	100.0	282	4	US-09-667-857-393	Sequence 393, Appl
3	1431	100.0	309	4	US-09-404-879A-392	Sequence 392, Appl
4	1431	100.0	309	4	US-09-667-857-392	Sequence 392, Appl
5	348	24.3	65	4	US-09-667-857-415	Sequence 415, Appl
6	246.5	17.2	316	4	US-09-910-174B-24	Sequence 24, Appl
7	246.5	17.2	316	4	US-09-620-461-24	Sequence 24, Appl
8	245	17.1	340	4	US-09-651-200-2	Sequence 2, Appl
9	245	17.1	441	4	US-09-651-200-4	Sequence 4, Appl
10	245	17.1	534	4	US-09-651-200-6	Sequence 6, Appl
11	245	17.1	534	4	US-09-651-200-24	Sequence 24, Appl
12	238.5	16.7	315	4	US-09-910-174B-28	Sequence 28, Appl
13	238.5	16.7	315	4	US-09-620-461-28	Sequence 28, Appl
14	223	15.6	513	4	US-09-910-174B-18	Sequence 18, Appl
15	223	15.6	513	4	US-09-620-461-18	Sequence 18, Appl
16	217.5	15.2	540	2	US-08-724-394A-4	Sequence 4, Appl
17	215.5	15.1	731	4	US-09-910-174B-15	Sequence 15, Appl
18	215.5	15.1	731	4	US-09-620-461-15	Sequence 15, Appl
19	213.5	14.9	584	4	US-09-910-174B-16	Sequence 16, Appl
20	213.5	14.9	584	4	US-09-620-461-16	Sequence 16, Appl
21	212.5	14.8	610	2	US-08-724-394A-5	Sequence 5, Appl
22	211.5	14.8	526	4	US-09-910-174B-9	Sequence 9, Appl
23	211.5	14.8	526	4	US-09-620-461-9	Sequence 9, Appl
24	211.5	14.8	526	4	US-09-949-016-6122	Sequence 6122, Ap
25	211.5	14.8	540	4	US-09-949-016-11644	Sequence 11644, A
26	211.5	14.8	589	2	US-08-724-394A-1	Sequence 1, Appl
27	207.5	14.5	119	4	US-09-910-174B-12	Sequence 12, Appl

; GENERAL INFORMATION:
 ; APPLICANT: Mitcham, Jennifer L.
 ; APPLICANT: King, Gordon E.
 ; APPLICANT: Algate, Paul A.
 ; APPLICANT: Fling, Steven P.
 ; APPLICANT: Retter, Marc W.
 ; APPLICANT: Panger, Gary Richard
 ; APPLICANT: Reed, Steven G.
 ; APPLICANT: Vedvick, Thomas S.
 ; APPLICANT: Carter, Darrick
 ; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
 ; TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
 ; FILE REFERENCE: 210121.462C5
 ; CURRENT APPLICATION NUMBER: US/09/667,857
 ; CURRENT FILING DATE: 2000-09-20
 ; NUMBER OF SEQ ID NOS: 455
 ; SOFTWARE: FastSeq for Windows Version 3.0
 ; SEQ ID NO 393
 ; LENGTH: 282
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 ; US-09-667-857-393

 Query Match 100.0%; Score 1431; DB 4; Length 282;
 Best Local Similarity 100.0%; Pred. No. 2.5e-138; Indels 0; Gaps 0;
 Matches 282; Conservative 0; Mismatches 0;

Qy 1 MASLGQILFWSIIIIIIILAGAIALLIIFGIGSRHSITVTTVASAGNIGEDGILSCTFEP 60
 Db 1 MASLGQILFWSIIIIIIILAGAIALLIIFGIGSRHSITVTTVASAGNIGEDGILSCTFEP 60
 Qy 61 DIKLSDIVIOWLKEGVLGLVHEPKEGKDELSEODEMFRGTAVFADQVIVGNASRLKNV 120
 Db 61 DIKLSDIVIOWLKEGVLGLVHEPKEGKDELSEODEMFRGTAVFADQVIVGNASRLKNV 120
 Qy 121 QLTDAQTYKCYIITSKGGNANLEYKTGAFSPMEVNVVDYNASSETLRCEAPRFPQPTVV 180
 Db 121 QLTDAQTYKCYIITSKGGNANLEYKTGAFSPMEVNVVDYNASSETLRCEAPRFPQPTVV 180
 Qy 181 WASQVDQGANFSEVSNFSFELNSNTVMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
 Db 181 WASQVDQGANFSEVSNFSFELNSNTVMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
 Qy 241 TESIKRSHQLLNKSKASLCVSSFFAISWALLPLSPYMLK 282
 Db 241 TESIKRSHQLLNKSKASLCVSSFFAISWALLPLSPYMLK 282

RESULT 3
 US-09-404-879A-392
 ; Sequence 392, Application US/09404879A
 ; Patent No. 6468546
 ; GENERAL INFORMATION:
 ; APPLICANT: Mitcham, Jennifer L.
 ; APPLICANT: King, Gordon E.
 ; APPLICANT: Algate, Paul A.
 ; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
 ; TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
 ; FILE REFERENCE: 210121.462C2
 ; CURRENT APPLICATION NUMBER: US/09/404,879A
 ; CURRENT FILING DATE: 1999-09-24
 ; NUMBER OF SEQ ID NOS: 393
 ; SOFTWARE: FastSeq for Windows Version 3.0
 ; SEQ ID NO 392
 ; LENGTH: 309
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 ; US-09-404-879A-392

 Query Match 100.0%; Score 1431; DB 4; Length 309;
 Best Local Similarity 100.0%; Pred. No. 2.8e-138; Indels 0; Gaps 0;
 Matches 282; Conservative 0; Mismatches 0;

Qy 1 MASLGQILFWSIIIIIIILAGAIALLIIFGIGSRHSITVTTVASAGNIGEDGILSCTFEP 60
 Db 1 MASLGQILFWSIIIIIIILAGAIALLIIFGIGSRHSITVTTVASAGNIGEDGILSCTFEP 60
 Qy 61 DIKLSDIVIOWLKEGVLGLVHEPKEGKDELSEODEMFRGTAVFADQVIVGNASRLKNV 120
 Db 61 DIKLSDIVIOWLKEGVLGLVHEPKEGKDELSEODEMFRGTAVFADQVIVGNASRLKNV 120
 Qy 121 QLTDAQTYKCYIITSKGGNANLEYKTGAFSPMEVNVVDYNASSETLRCEAPRFPQPTVV 180
 Db 121 QLTDAQTYKCYIITSKGGNANLEYKTGAFSPMEVNVVDYNASSETLRCEAPRFPQPTVV 180
 Qy 181 WASQVDQGANFSEVSNFSFELNSNTVMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
 Db 181 WASQVDQGANFSEVSNFSFELNSNTVMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
 Qy 241 TESIKRSHQLLNKSKASLCVSSFFAISWALLPLSPYMLK 282
 Db 241 TESIKRSHQLLNKSKASLCVSSFFAISWALLPLSPYMLK 282

Qy 1 MASLGQILFWSIIIIIIILAGAIALLIIFGIGSRHSITVTTVASAGNIGEDGILSCTFEP 60
 Db 28 MASLGQILFWSIIIIIIILAGAIALLIIFGIGSRHSITVTTVASAGNIGEDGILSCTFEP 87
 Qy 61 DIKLSDIVIOWLKEGVLGLVHEPKEGKDELSEODEMFRGTAVFADQVIVGNASRLKNV 120
 Db 88 DIKLSDIVIOWLKEGVLGLVHEPKEGKDELSEODEMFRGTAVFADQVIVGNASRLKNV 147
 Qy 121 QLTDAQTYKCYIITSKGGNANLEYKTGAFSPMEVNVVDYNASSETLRCEAPRFPQPTVV 180
 Db 148 QLTDAQTYKCYIITSKGGNANLEYKTGAFSPMEVNVVDYNASSETLRCEAPRFPQPTVV 207
 Qy 181 WASQVDQGANFSEVSNFSFELNSNTVMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
 Db 208 WASQVDQGANFSEVSNFSFELNSNTVMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 267
 Qy 241 TESIKRSHQLLNKSKASLCVSSFFAISWALLPLSPYMLK 282
 Db 268 TESIKRSHQLLNKSKASLCVSSFFAISWALLPLSPYMLK 309

RESULT 4
 US-09-667-857-392
 ; Sequence 392, Application US/09667857
 ; Patent No. 6699664
 ; GENERAL INFORMATION:
 ; APPLICANT: Mitcham, Jennifer L.
 ; APPLICANT: King, Gordon E.
 ; APPLICANT: Algate, Paul A.
 ; APPLICANT: Fling, Steven P.
 ; APPLICANT: Retter, Marc W.
 ; APPLICANT: Panger, Gary Richard
 ; APPLICANT: Reed, Steven G.
 ; APPLICANT: Vedvick, Thomas S.
 ; APPLICANT: Carter, Darrick
 ; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
 ; TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
 ; FILE REFERENCE: 210121.462C5
 ; CURRENT APPLICATION NUMBER: US/09/667,857
 ; CURRENT FILING DATE: 2000-09-20
 ; NUMBER OF SEQ ID NOS: 455
 ; SOFTWARE: FastSeq for Windows Version 3.0
 ; SEQ ID NO 392
 ; LENGTH: 309
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 ; US-09-667-857-392

 Query Match 100.0%; Score 1431; DB 4; Length 309;
 Best Local Similarity 100.0%; Pred. No. 2.8e-138; Indels 0; Gaps 0;
 Matches 282; Conservative 0; Mismatches 0;

Qy 1 MASLGQILFWSIIIIIIILAGAIALLIIFGIGSRHSITVTTVASAGNIGEDGILSCTFEP 60
 Db 28 MASLGQILFWSIIIIIIILAGAIALLIIFGIGSRHSITVTTVASAGNIGEDGILSCTFEP 87
 Qy 61 DIKLSDIVIOWLKEGVLGLVHEPKEGKDELSEODEMFRGTAVFADQVIVGNASRLKNV 120
 Db 88 DIKLSDIVIOWLKEGVLGLVHEPKEGKDELSEODEMFRGTAVFADQVIVGNASRLKNV 147
 Qy 121 QLTDAQTYKCYIITSKGGNANLEYKTGAFSPMEVNVVDYNASSETLRCEAPRFPQPTVV 180
 Db 148 QLTDAQTYKCYIITSKGGNANLEYKTGAFSPMEVNVVDYNASSETLRCEAPRFPQPTVV 207
 Qy 181 WASQVDQGANFSEVSNFSFELNSNTVMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
 Db 208 WASQVDQGANFSEVSNFSFELNSNTVMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 267
 Qy 241 TESIKRSHQLLNKSKASLCVSSFFAISWALLPLSPYMLK 282
 Db 268 TESIKRSHQLLNKSKASLCVSSFFAISWALLPLSPYMLK 309

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RESULT 5
US-09-667-857-415
; Sequence 415, Application US/09667857
; Patent No. 6699664
; GENERAL INFORMATION:
; APPLICANT: Mitcham, Jennifer L.
; APPLICANT: King, Gordon E.
; APPLICANT: Algate, Paul A.
; APPLICANT: Fling, Steven P.
; APPLICANT: Retter, Marc W.
; APPLICANT: Fanger, Gary Richard
; APPLICANT: Reed, Steven G.
; APPLICANT: Vedvick, Thomas S.
; APPLICANT: Carter, Barrick
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
; TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
; FILE REFERENCE: 210121.462C5
; CURRENT APPLICATION NUMBER: US/09/667,857
; CURRENT FILING DATE: 2000-09-20
; NUMBER OF SEQ ID NOS: 455
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 415
; LENGTH: 65
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-667-857-415

Query Match      24.3%; Score 348; DB 4; Length 65;
Best Local Similarity 100.0%; Pred. No. 2.9e-28;
Matches 65; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 136 KGKGNANLEYKTGA-FSMPEVNVNDYNASSETLRCEAPRWFPOPTVWASQVDQGANFSEVS 195
Db 1 KGKGNANLEYKTGA-FSMPEVNVNDYNASSETLRCEAPRWFPOPTVWASQVDQGANFSEVS 60

Qy 196 NTSFE 200
Db 61 NTSFE 65

RESULT 6
US-09-910-174B-24
; Sequence 24, Application US/09910174B
; Patent No. 6630575
; GENERAL INFORMATION:
; APPLICANT: Coyle, Anthony J.
; APPLICANT: Fraser, Christopher C.
; APPLICANT: Manning, Stephen
; TITLE OF INVENTION: B7-H2 Molecules, No. 6630575el Members of the B7
; TITLE OF INVENTION: Family and Uses Thereof
; FILE REFERENCE: 35800/236924
; CURRENT APPLICATION NUMBER: US/09/910,174B
; CURRENT FILING DATE: 2001-07-20
; PRIOR APPLICATION NUMBER: US 09/620,461
; PRIOR FILING DATE: 2000-07-20
; NUMBER OF SEQ ID NOS: 32
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 24
; LENGTH: 316
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-910-174B-24

Query Match      17.2%; Score 246.5; DB 4; Length 316;
Best Local Similarity 30.2%; Pred. No. 9e-17;
Matches 70; Conservative 44; Mismatches 99; Indels 19; Gaps 9;

Qy 21 GATALIIGFGISGRHSITVTTVASAGNIGEDGILSCTF--EPDIKLSDIVIQWLKEGVLG 78
Db 15 GAALGALWFCLTGALEVOQVPDPVVALVGTDTATLCCSFSPSPGSLAQLNLWQLTDTKQ 74

Qy 79 LVHEFKEGKDELSQDEMFRGRTAVFADQVTVGNASRLKNVQLTDAQTYKCYIITSKKG 138
Db 75 LVHSAFEGQD---QGSAYANRRTALFPDLLAQGNASRLQVRVADEGSFTCF-VSIRDF 129

Qy 139 GNaNLEYKTGA-FSMPEVNVNDYN-----ASSETLRCEAPRWFPOPTVWASQVDQGANFS 192
Db 130 GSAAVSLQVAAPYKPSMTLEPNKDLRPGDVTITCSSYRGYPAEVFW--QDGGVPLT 187

Qy 193 EVSNTSPELSENVTMKVSVLYNVT-INNTYSCHIENDIAK--ATGDIKVT 241
Db 188 GNVTTT-QMANEQGLFDVHSLRVVLGANGTYSCLVRNPVLQDQAHGSVTIT 238

RESULT 7
US-09-620-461-24
; Sequence 24, Application US/09620461
; Patent No. 6635750
; GENERAL INFORMATION:
; APPLICANT: Coyle, Anthony J.
; APPLICANT: Fraser, Christopher C.
; APPLICANT: Manning, Stephen
; TITLE OF INVENTION: B7-H2 Molecules, No. 6635750el Members of the B7
; TITLE OF INVENTION: Family and Uses Thereof
; FILE REFERENCE: 5800-149
; CURRENT APPLICATION NUMBER: US/09/620,461
; CURRENT FILING DATE: 2000-07-20
; NUMBER OF SEQ ID NOS: 29
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 24
; LENGTH: 316
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-620-461-24

Query Match      17.2%; Score 246.5; DB 4; Length 316;
Best Local Similarity 30.2%; Pred. No. 9e-17;
Matches 70; Conservative 44; Mismatches 99; Indels 19; Gaps 9;

Qy 21 GATALIIGFGISGRHSITVTTVASAGNIGEDGILSCTF--EPDIKLSDIVIQWLKEGVLG 78
Db 15 GAALGALWFCLTGALEVOQVPDPVVALVGTDTATLCCSFSPSPGSLAQLNLWQLTDTKQ 74

Qy 79 LVHEFKEGKDELSQDEMFRGRTAVFADQVTVGNASRLKNVQLTDAQTYKCYIITSKKG 138
Db 75 LVHSAFEGQD---QGSAYANRRTALFPDLLAQGNASRLQVRVADEGSFTCF-VSIRDF 129

Qy 139 GNaNLEYKTGA-FSMPEVNVNDYN-----ASSETLRCEAPRWFPOPTVWASQVDQGANFS 192
Db 130 GSAAVSLQVAAPYKPSMTLEPNKDLRPGDVTITCSSYRGYPAEVFW--QDGGVPLT 187

Qy 193 EVSNTSPELSENVTMKVSVLYNVT-INNTYSCHIENDIAK--ATGDIKVT 241
Db 188 GNVTTT-QMANEQGLFDVHSLRVVLGANGTYSCLVRNPVLQDQAHGSVTIT 238

RESULT 8
US-09-651-200-2
; Sequence 2, Application US/09651200
; Patent No. 6429303
; GENERAL INFORMATION:
; APPLICANT: Green et al
; TITLE OF INVENTION: Polynucleotides Encoding Members of the Human B
; TITLE OF INVENTION: Lymphocyte Activation Antigen B-7 Family and
; TITLE OF INVENTION: Polypeptides Encoded Thereby
; FILE REFERENCE: 15966-562 (CURA-62)
; CURRENT APPLICATION NUMBER: US/09/651,200
; CURRENT FILING DATE: 2000-08-30
; PRIOR APPLICATION NUMBER: 60/152383
; PRIOR FILING DATE: 1999-09-03
; PRIOR APPLICATION NUMBER: 60/172909
; PRIOR FILING DATE: 1999-12-21
; PRIOR APPLICATION NUMBER: 60/183578
; PRIOR FILING DATE: 2000-02-18
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: Patent in Ver. 2.0
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166 LRCEAPRWFPQPTVVMASQVDQGANFSEVSNTPFELNSENVTMKVSVLYNVT-INNTYS 224
; SEQ ID NO 2
; LENGTH: 340
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-651-200-2

Query Match 17.1%; Score 245; DB 4; Length 340;
Best Local Similarity 27.8%; Pred. No. 1.4e-16;
Matches 72; Conservative 50; Mismatches 101; Indels 36; Gaps 11;

Qy 11 SIISIIILAGAILIIGFI---SGRHSITVTTVAS-----AGNIGEDGI 53
Db 12 SILRVVLGANGTYSCLVRNPVLQDDAHSSVTIIPQSPGTGAVEVQVPDPVVALVGTDTAT 71

Qy 54 LSCTF--EPDIKLSDIVIOMLKEGVGLVHEFKGKDELSEQDEMFRGTAVFADQVIVG 111
Db 72 LHCFSFEPGFSITQLNLIWQLTDTKQLVHSFTEGRD---QGSAYANRTALFPDIIAQQ 127

Qy 112 NASLRKNVOLTADGTYKCVIITSKGNANLEYKTGA-FSMPEVNVN---ASSET 165
Db 128 NASLRQVRVADEGSFTCF-VSIRDFGSAVSLQVAAPYKPSMTLEPNKDLRPGDVT 186

Qy 166 LRCEAPRWFPQPTVVMASQVDQGANFSEVSNTPFELNSENVTMKVSVLYNVT-INNTYS 224
Db 187 ITCSYRGYPEAEVFW--QDQGVPLTGNVTTS-QMANEQGLFDVHSLRVVLGANGTYS 243

Qy 225 CMIENDIAK--ATGDIKVT 241
Db 244 CLVRNPVLQDDAHGVSVTIT 262

RESULT 9
US-09-651-200-4
; Sequence 4, Application US/09651200
; Patent No. 6429303
; GENERAL INFORMATION:
; APPLICANT: Green et al
; TITLE OF INVENTION: Polynucleotides Encoding Members of the Human B
; TITLE OF INVENTION: Lymphocyte Activation Antigen B-7 Family and
; TITLE OF INVENTION: Polypeptides Encoded Thereby
; FILE REFERENCE: 15966-562 (CURA-62)
; CURRENT APPLICATION NUMBER: US/09/651,200
; CURRENT FILING DATE: 2000-08-30
; PRIOR APPLICATION NUMBER: 60/152383
; PRIOR FILING DATE: 1999-09-03
; PRIOR APPLICATION NUMBER: 60/172909
; PRIOR FILING DATE: 1999-12-21
; PRIOR APPLICATION NUMBER: 60/183578
; PRIOR FILING DATE: 2000-02-18
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 4
; LENGTH: 441
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-651-200-4

Query Match 17.1%; Score 245; DB 4; Length 441;
Best Local Similarity 27.8%; Pred. No. 2.2e-16;
Matches 72; Conservative 50; Mismatches 101; Indels 36; Gaps 11;

Qy 11 SIISIIILAGAILIIGFI---SGRHSITVTTVAS-----AGNIGEDGI 53
Db 113 SILRVVLGANGTYSCLVRNPVLQDDAHSSVTIIPQSPGTGAVEVQVPDPVVALVGTDTAT 172

Qy 54 LSCTF--EPDIKLSDIVIOMLKEGVGLVHEFKGKDELSEQDEMFRGTAVFADQVIVG 111
Db 173 LHCFSFEPGFSITQLNLIWQLTDTKQLVHSFTEGRD---QGSAYANRTALFPDIIAQQ 228

Qy 112 NASLRKNVOLTADGTYKCVIITSKGNANLEYKTGA-FSMPEVNVN---ASSET 165
Db 229 NASLRQVRVADEGSFTCF-VSIRDFGSAVSLQVAAPYKPSMTLEPNKDLRPGDVT 287

166 LRCEAPRWFPQPTVVMASQVDQGANFSEVSNTPFELNSENVTMKVSVLYNVT-INNTYS 224
288 ITCSYRGYPEAEVFW--QDQGVPLTGNVTTS-QMANEQGLFDVHSLRVVLGANGTYS 344
225 CMIENDIAK--ATGDIKVT 241
345 CLVRNPVLQDDAHGVSVTIT 363

RESULT 10
US-09-651-200-6
; Sequence 6, Application US/09651200
; Patent No. 6429303
; GENERAL INFORMATION:
; APPLICANT: Green et al
; TITLE OF INVENTION: Polynucleotides Encoding Members of the Human B
; TITLE OF INVENTION: Lymphocyte Activation Antigen B-7 Family and
; TITLE OF INVENTION: Polypeptides Encoded Thereby
; FILE REFERENCE: 15966-562 (CURA-62)
; CURRENT APPLICATION NUMBER: US/09/651,200
; CURRENT FILING DATE: 2000-08-30
; PRIOR APPLICATION NUMBER: 60/152383
; PRIOR FILING DATE: 1999-09-03
; PRIOR APPLICATION NUMBER: 60/172909
; PRIOR FILING DATE: 1999-12-21
; PRIOR APPLICATION NUMBER: 60/183578
; PRIOR FILING DATE: 2000-02-18
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 6
; LENGTH: 534
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-651-200-6

Query Match 17.1%; Score 245; DB 4; Length 534;
Best Local Similarity 27.8%; Pred. No. 2.9e-16;
Matches 72; Conservative 50; Mismatches 101; Indels 36; Gaps 11;

Qy 11 SIISIIILAGAILIIGFI---SGRHSITVTTVAS-----AGNIGEDGI 53
Db 206 SILRVVLGANGTYSCLVRNPVLQDDAHSSVTIIPQSPGTGAVEVQVPDPVVALVGTDTAT 265

Qy 54 LSCTF--EPDIKLSDIVIOMLKEGVGLVHEFKGKDELSEQDEMFRGTAVFADQVIVG 111
Db 266 LRCFSFEPGFSITQLNLIWQLTDTKQLVHSFTEGRD---QGSAYANRTALFPDIIAQQ 321

Qy 112 NASLRKNVOLTADGTYKCVIITSKGNANLEYKTGA-FSMPEVNVN---ASSET 165
Db 322 NASLRQVRVADEGSFTCF-VSIRDFGSAVSLQVAAPYKPSMTLEPNKDLRPGDVT 380

Qy 166 LRCEAPRWFPQPTVVMASQVDQGANFSEVSNTPFELNSENVTMKVSVLYNVT-INNTYS 224
Db 381 ITCSYRGYPEAEVFW--QDQGVPLTGNVTTS-QMANEQGLFDVHSLRVVLGANGTYS 437

Qy 225 CMIENDIAK--ATGDIKVT 241
Db 438 CLVRNPVLQDDAHGVSVTIT 456

RESULT 11
US-09-651-200-24
; Sequence 24, Application US/09651200
; Patent No. 6429303
; GENERAL INFORMATION:
; APPLICANT: Green et al
; TITLE OF INVENTION: Polynucleotides Encoding Members of the Human B
; TITLE OF INVENTION: Lymphocyte Activation Antigen B-7 Family and
; TITLE OF INVENTION: Polypeptides Encoded Thereby
; FILE REFERENCE: 15966-562 (CURA-62)
; CURRENT APPLICATION NUMBER: US/09/651,200
; CURRENT FILING DATE: 2000-08-30
; PRIOR APPLICATION NUMBER: 60/152383

;; CURRENT APPLICATION NUMBER: US/09/910,174B
;; CURRENT FILING DATE: 2001-07-20
;; PRIOR APPLICATION NUMBER: US 09/620,461
;; PRIOR FILING DATE: 2000-07-20
;; NUMBER OF SEQ ID NOS: 32
;; SOFTWARE: FastSEQ for Windows Version 4.0
;; SEQ ID NO 18
;; LENGTH: 513
;; TYPE: PRT
;; ORGANISM: Homo sapiens
US-09-910-174B-18

Query Match 15.6%; Score 223; DB 4; Length 513;
Best Local Similarity 26.4%; Pred. No. 5e-14;
Matches 69; Conservative 44; Mismatches 108; Indels 40; Gaps 10;
QY 35 HSIITVTVASAGNI---GEGILSCTFEPDIKLSDIVIOWLKEGVLGVHFEKKGDEL 90
DB 27 HSAQFVLGSPGPIAMVGEDADLPCHLPTMSAETMELKWSVSSLRQVNVVYADGKEVE 86
QY 91 SEQDEMFRGRTAVFADQVIVGNASLRKKNVQLTDAGTYKCYIITSKGGNANLEYKTGAF 150
DB 87 DRQSAFYRGRTSILRDGITAGKALRIHNVTASDSKYLCTYFQDGFYKALVELKVAAL 146
QY 151 SMPENVVD---YNASSETLRCEAPRWFPQPTVVWASQVDQGANFSEVNTSFELNSENV 207
DB 147 G-SDLHVDVKYKGGIHLKCRSTGWYPOQIQWSN--NKGEN---IPTVEAPVVADGVG 200
QY 208 MKVY--SVLYNVNTINNTYSCMIENDIAKATGDIKVTESEIKRRSHLQLLNSKASLCVSS- 264
DB 201 LYAFAASVIMRGSGEGVSCIT-----RSSLLGLEKTASISIAADP 240
QY 265 FF--AISW--ALLPLSPYML 281
DB 241 FFRSAQRWIAALARTLPVLL 261

RESULT 15
US-09-620-461-18
;; Sequence 18, Application US/09620461
;; Patent No. 6635750
;; GENERAL INFORMATION:
;; APPLICANT: Coyle, Anthony J.
;; APPLICANT: Fraser, Christopher C.
;; APPLICANT: Manning, Stephen
;; TITLE OF INVENTION: B7-H2 Molecules, No. 6635750el Members of the B7
;; FILE OF INVENTION: Family and Uses Thereof
;; FILE REFERENCE: 5800-149
;; CURRENT APPLICATION NUMBER: US/09/620,461
;; CURRENT FILING DATE: 2000-07-20
;; NUMBER OF SEQ ID NOS: 29
;; SOFTWARE: FastSEQ for Windows Version 3.0
;; SEQ ID NO 18
;; LENGTH: 513
;; TYPE: PRT
;; ORGANISM: Homo sapiens
US-09-620-461-18

Query Match 15.6%; Score 223; DB 4; Length 513;
Best Local Similarity 26.4%; Pred. No. 5e-14;
Matches 69; Conservative 44; Mismatches 108; Indels 40; Gaps 10;
QY 35 HSIITVTVASAGNI---GEGILSCTFEPDIKLSDIVIOWLKEGVLGVHFEKKGDEL 90
DB 27 HSAQFVLGSPGPIAMVGEDADLPCHLPTMSAETMELKWSVSSLRQVNVVYADGKEVE 86
QY 91 SEQDEMFRGRTAVFADQVIVGNASLRKKNVQLTDAGTYKCYIITSKGGNANLEYKTGAF 150
DB 87 DRQSAFYRGRTSILRDGITAGKALRIHNVTASDSKYLCTYFQDGFYKALVELKVAAL 146
QY 151 SMPENVVD---YNASSETLRCEAPRWFPQPTVVWASQVDQGANFSEVNTSFELNSENV 207
DB 147 G-SDLHVDVKYKGGIHLKCRSTGWYPOQIQWSN--NKGEN---IPTVEAPVVADGVG 200

QY 208 MKVY--SVLYNVNTINNTYSCMIENDIAKATGDIKVTESEIKRRSHLQLLNSKASLCVSS- 264
DB 201 LYAFAASVIMRGSGEGVSCIT-----RSSLLGLEKTASISIAADP 240
QY 265 FF--AISW--ALLPLSPYML 281
DB 241 FFRSAQRWIAALARTLPVLL 261
Search completed: April 19, 2005, 07:17:34
Job time : 30 secs

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: April 19, 2005, 04:13:09 ; Search time 85 Seconds
(without alignments)
1698.898 Million cell updates/sec

Title: US-10-773-715-6
Perfect score: 1431
Sequence: 1 MASLGQLFWSIIIIIIIIA.....SSPFAISWALLPLSPYMLK 282

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1612378 seqs, 512079187 residues

Total number of hits satisfying chosen parameters: 1612378

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : UniProt_03.*
1: uniprot_sprot.*
2: uniprot_trembl.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	1431	100.0	282	2 Q7Z7D3	Q7Z7D3 homo sapien
2	1425	99.6	282	2 Q9H6B2	Q9H6B2 homo sapien
3	1261.5	88.2	283	2 Q7TPH5	Q7TPH5 mus musculus
4	1257.5	87.9	283	2 Q7TSP5	Q7TSP5 mus musculus
5	1253.5	87.6	283	2 Q8K091	Q8K091 mus musculus
6	958	66.9	187	2 Q6P037	Q6P037 homo sapien
7	512	35.8	285	2 Q7ZY30	Q7ZY30 xenopus lae
8	493.5	34.5	276	2 Q640S5	Q640S5 xenopus tro
9	448	31.3	275	2 Q8AVV1	Q8AVV1 xenopus lae
10	276	19.3	465	2 Q640J0	Q640J0 xenopus lae
11	255.5	17.9	305	2 Q6DJ75	Q6DJ75 xenopus tro
12	246.5	17.2	316	2 Q9BXR1	Q9BXR1 homo sapien
13	245	17.1	388	2 Q8NC34	Q8NC34 homo sapien
14	245	17.1	493	2 Q6PSY4	Q6PSY4 homo sapien
15	245	17.1	533	2 Q8NCB6	Q8NCB6 homo sapien
16	245	17.1	534	2 Q8NB18	Q8NB18 homo sapien
17	240.5	16.8	316	2 Q6UX12	Q6UX12 homo sapien
18	235	16.4	316	2 Q8VE38	Q8VE38 mus musculus
19	232	16.2	316	2 Q7TPB4	Q7TPB4 rattus norv
20	225.5	15.8	466	2 Q6UXE8	Q6UXE8 homo sapien
21	225	15.7	220	2 Q9NM06	Q9NM06 homo sapien
22	225	15.7	414	2 Q9UM44	Q9UM44 homo sapien
23	223	15.6	495	2 Q9HCY1	Q9HCY1 homo sapien
24	223	15.6	513	2 Q00481	Q00481 homo sapien
25	222.5	15.5	347	2 Q9H730	Q9H730 homo sapien
26	222.5	15.5	500	2 Q6UX41	Q6UX41 homo sapien
27	219.5	15.3	286	2 Q46535	Q46535 bos taurus
28	217	15.2	280	2 Q73716	Q73716 grus americ
29	215.5	15.1	304	2 Q9BE26	Q9BE26 macaca fasc
30	215.5	15.1	526	1 BUTY BOVIN	P18892 bos taurus
31	215.5	15.1	731	2 P78409	P78409 homo sapien

RESULT 1

ID	Q7Z7D3	PRELIMINARY;	PRT;	282 AA.
AC	Q7Z7D3:			
DT	01-OCT-2003	(TrEMBLrel. 25, Created)		
DT	01-OCT-2003	(TrEMBLrel. 25, Last sequence update)		
DT	25-OCT-2004	(TrEMBLrel. 28, Last annotation update)		
DE	Immune costimulatory protein B7-H4 (T cell costimulatory molecule B7x) (B7h.5).			
DE	Name=B7-H4; ORFNames=UNQ659;			
GN	Homo sapiens (Human)			
OS	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Rutheria; Primates; Catarrhini; Hominidae; Homo.			
OC	NCBI_TaxID=9606;			
OX	[1]			
RN	SEQUENCE FROM N.A.			
RP	Sica G.L., Choi I.-H., Zhu G., Tamada K., Wang S.-D., Tamura H., Chapoval A.I., Flies D.B., Bajorath J., Chen L.; Submitted (APR-2003) to the EMBL/GenBank/DBJ databases.			
RA	[2]			
RN	SEQUENCE FROM N.A.			
RP	MEDLINE=22833980; PubMed=12920180; DOI=10.1073/pnas.1434299100; Zang X., Loke P., Kim J., Murphy K., Waitz R., Allison J.P.;			
RA	"B7x: a widely expressed B7 family member that inhibits T cell activation.";			
RT	Proc. Natl. Acad. Sci. U.S.A. 100:10388-10392(2003).			
RL	[3]			
RN	SEQUENCE FROM N.A.			
RP	MEDLINE=22887296; PubMed=12975309; DOI=10.1101/gr.1293003; Clark H.F., Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J., Chen J., Chow B., Chui C., Crowley C., Currell B., Deuel B., Dowd P., Eaton D., Foster J., Grimaldi C., Gu Q., Hass P.E., Heldens S., Huang A., Kim H.S., Klimowski L., Jin Y., Johnson S., Lee J., Lewis L., Liao D., Mark M., Robbie E., Sanchez C., Schoenfeld J., Seehagiri S., Simmons L., Singh J., Smith V., Stinson J., Vagts A., Vandlen R., Watanabe C., Wieand D., Woods K., Xie M.H., Yansura D., Yi S., Yu G., Yuan J., Zhang M., Zhang Z., Goddard A., Wood W.I., Godowski P.;			
RA	"The secreted protein discovery initiative (SPDI), a large-scale effort to identify novel human secreted and transmembrane proteins: a bioinformatics assessment.";			
RT	Bioinformatics assessment.";			
RL	Genome Res. 13:2265-2270(2003).			
RP	[4]			
RN	SEQUENCE FROM N.A.			
RP	TISSUE=Brain;			
RC	MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;			
RX	Straussberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D., Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K., Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Haie F., Diatchenko L., Marusina K., Farmer A., Rubin G.M., Hong L., Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E., Brownstein M.J., Udén T.B., Toshiyuki S., Carninci P., Prange C., Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,			

ALIGNMENTS

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RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahy J., Helton E., Kettelman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krywinski M.I., Skalski U., Smailus D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Marra M.A.;
RA "Generation and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences.",
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN [5]
RN TISSUE=Brain;
RC Director MGC Project;
RL Submitted (JUN-2004) to the EMBL/GenBank/DBJ databases.
DR EMBL; AY280972; AAP37283.1; -
DR EMBL; AY346100; AAQ24206.1; -
DR EMBL; AY358352; AAQ88718.1; -
DR EMBL; BC074729; AAH74729.1; -
DR HSSP; O63345; 1PKO.
DR InterPro; IPR007110; Ig-like.
DR Pfam; PF00047; Ig; 1. LIKE; 2.
DR PROSITE; PS50835; IG_LIKE; 2.
SQ SEQUENCE 282 AA; 30878 MW; 1C9C565A9242E78C CRC64;

Query Match 100.0%; Score 1431; DB 2; Length 282;
Best Local Similarity 100.0%; Pred. No. 1.5e-107;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MASLGQILFWSIISIIIIAGAIALLIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60
DB 1 MASLGQILFWSIISIIIIAGAIALLIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60

QY 61 DIKLSDIVIOWLKEGVLGVHFEKKGDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
DB 61 DIKLSDIVIOWLKEGVLGVHFEKKGDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120

QY 121 QLTDACTYKCYIITSKGKNANLEYKTGAFSPMPVNVNDYNASSETLRCEAPRFPQPTVV 180
DB 121 QLTDACTYKCYIITSKGKNANLEYKTGAFSPMPVNVNDYNASSETLRCEAPRFPQPTVV 180

QY 181 WASQVDQGANFSEVNTSFELNSENVTMKVSVLVNNTYSCMIENDIAKATGDIKV 240
DB 181 WASQVDQGANFSEVNTSFELNSENVTMKVSVLVNNTYSCMIENDIAKATGDIKV 240

QY 241 TESEIKRRSHLQLLNKSKASLCVSSPFAISWALLPLSPYMLK 282
DB 241 TESEIKRRSHLQLLNKSKASLCVSSPFAISWALLPLSPYMLK 282

RESULT 2
QY Q9H6B2 PRELIMINARY; PRT; 282 AA.
ID Q9H6B2
AC Q9H6B2;
DT 01-MAR-2001 (Tremblrel. 16, Created)
DT 01-MAR-2001 (Tremblrel. 16, Last sequence update)
DT 01-OCT-2003 (Tremblrel. 25, Last annotation update)
DE Hypothetical protein FLJ22418.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RN SEQUENCE FROM N.A.
RA Kawabata A., Hikiji T., Kobatake N., Inagaki H., Ikema Y., Okamoto S.,
RA Okitani R., Ota T., Suzuki Y., Obayashi M., Nishi T., Shibahara T.,
RA Tanaka T., Nakamura Y., Isoigai T., Sugano S.;
RL Submitted (AUG-2000) to the EMBL/GenBank/DBJ databases.
DR EMBL; AK026071; BAB15349.1; -
DR HSSP; O63345; 1PKO.
DR InterPro; IPR003599; Ig.
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DR InterPro; IPR007110; Ig-like.
DR SMART; SM00409; IG; 1.
DR PROSITE; PS50835; IG_LIKE; 2.
SQ SEQUENCE 282 AA; 30893 MW; 6F9066999A1EBDB4 CRC64;

Query Match 99.6%; Score 1425; DB 2; Length 282;
Best Local Similarity 99.6%; Pred. No. 4.7e-107;
Matches 281; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 MASLGQILFWSIISIIIIAGAIALLIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60
DB 1 MASLGQILFWSIISIIIIAGAIALLIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60

QY 61 DIKLSDIVIOWLKEGVLGVHFEKKGDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
DB 61 DIKLSDIVIOWLKEGVLGVHFEKKGDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120

QY 121 QLTDACTYKCYIITSKGKNANLEYKTGAFSPMPVNVNDYNASSETLRCEAPRFPQPTVV 180
DB 121 QLTDACTYKCYIITSKGKNANLEYKTGAFSPMPVNVNDYNASSETLRCEAPRFPQPTVV 180

QY 181 WASQVDQGANFSEVNTSFELNSENVTMKVSVLVNNTYSCMIENDIAKATGDIKV 240
DB 181 WASQVDQGANFSEVNTSFELNSENVTMKVSVLVNNTYSCMIENDIAKATGDIKV 240

QY 241 TESEIKRRSHLQLLNKSKASLCVSSPFAISWALLPLSPYMLK 282
DB 241 TESEIKRRSHLQLLNKSKASLCVSSPFAISWALLPLSPYMLK 282

RESULT 3
QY Q7TPH5 PRELIMINARY; PRT; 283 AA.
ID Q7TPH5
AC Q7TPH5;
DT 01-OCT-2003 (Tremblrel. 25, Created)
DT 01-OCT-2003 (Tremblrel. 25, Last sequence update)
DT 01-MAR-2004 (Tremblrel. 26, Last annotation update)
DE B7S1.
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Mus.
OX NCBI_TaxID=10090;
RN [1]
RN SEQUENCE FROM N.A.
RA MEDLINE=22703430; PubMed=12818166; DOI=10.1016/S1074-7613(03)00147-X;
RA Prasad D.V., Richards S., Mai X.M., Dong C.;
RA "B7S1, a novel B7 family member that negatively regulates T cell
RA activation.",
RA Immunity 18:863-873(2003).
RA EMBL; AY322147; AAP8965.1; -
RA HSSP; O63345; 1PKO.
RA InterPro; IPR003599; Ig.
RA InterPro; IPR007110; Ig-like.
RA SMART; SM00409; IG; 1.
RA PROSITE; PS50835; IG_LIKE; 2.
SQ SEQUENCE 283 AA; 30847 MW; A97F17461857850B CRC64;

Query Match 88.2%; Score 1261.5; DB 2; Length 283;
Best Local Similarity 88.0%; Pred. No. 8e-94;
Matches 249; Conservative 13; Mismatches 20; Indels 1; Gaps 1;

QY 1 MASLGQILFWSIISIIIIAGAIALLIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60
DB 1 MASLGQILFWSIISIIIIAGAIALLIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60

QY 61 DIKLSDIVIOWLKEGVLGVHFEKKGDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
DB 61 DIKLSDIVIOWLKEGVLGVHFEKKGDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120

QY 121 QLTDACTYKCYIITSKGKNANLEYKTGAFSPMPVNVNDYNASSETLRCEAPRFPQPTVV 180
DB 121 QLTDACTYKCYIITSKGKNANLEYKTGAFSPMPVNVNDYNASSETLRCEAPRFPQPTVV 180
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Qy 181 WASQVDOGANFSEVSNSTFELNSNVTKVSVLVYNTVNTTSCMIENDIAKATGDIKV 240
Db 181 WASQVDOGANFSEVSNSTFELNSNVTKVSVLVYNTVNTTSCMIENDIAKATGDIKV 240

Qy 241 TESIKRSHLQLNLSKASLCV-SSFFAISWALLPLSPYMLK 282
Db 241 TDSEVKRRSQQLNLSGSPCVFSSAFAGWALLSLSCCLMLR 283

RESULT 4
Q7TSP5
ID Q7TSP5 PRELIMINARY; PRT; 283 AA.
AC Q7TSP5,
DT 01-OCT-2003 (TrEMBLrel. 25, Created)
DT 01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
DT 05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
DE Immune costimulatory protein B7-H4 (T cell costimulatory molecule
DE B7x).
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=BALB/c;
RA Sica G.L., Choi I.-H., Zhu G., Tamada K., Wang S.-D., Tamura H.,
RA Chapoval A.I., Flies D.B., Bajorath J., Chen L.;
RL Submitted (APR-2003) to the EMBL/GenBank/DBJ databases.
RN [2]
RP SEQUENCE FROM N.A.
RC STRAIN=FVB/N;
RX MEDLINE=22833980; PubMed=12920180; DOI=10.1073/pnas.1434299100;
RA Zang X., Loke P., Kim J., Murphy K., Waitz R., Allison J.P.;
RT "B7x: a widely expressed B7 family member that inhibits T cell
RT activation."
RL Proc. Natl. Acad. Sci. U.S.A. 100:10388-10392(2003).
DR EMBL; AY280973; AAF37284.1; -
DR EMBL; AY346099; AAQ24205.1; -
DR HSP; Q63345; 1PKO.
DR InterPro; IPR007110; IG-like.
DR PROSITE; PS50835; IG LIKE; 2.
SQ SEQUENCE 283 AA; 30875 MW; 7E2F174618578519 CRC64;

Query Match 87.9%; Score 1257.5; DB 2; Length 283;
Best Local Similarity 87.6%; Pred. No. 1.7e-93;
Matches 248; Conservative 13; Mismatches 21; Indels 1; Gaps 1;

Qy 1 MASLGQLFWSIIIIILAGAIALLIIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP 60
Db 1 MASLGQIIFWSIINIILAGAIALLIIGFGISGRHFTVTTFTSAGNIGEDGILSCTFEP 60

Qy 61 DIKLSDIVIOWLKEGVLGVHEFKEGKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
Db 61 DIKLVGIVOWLKEGIVKGLVHEFKEGKDDLSQQHEMFRGRTAVFADQVIVGNASRLKNV 120

Qy 121 QLTDAGTYKCYIITSKGNANLEYKYGAFSMPVNVYDYNASSETLRCEAPRPFQPTVV 180
Db 121 QLTDAGTYCYIRTSTKGNANLEYKYGAFSMPVNVYDYNASSETLRCEAPRPFQPTVA 180

Qy 181 WASQVDOGANFSEVSNSTFELNSNVTKVSVLVYNTVNTTSCMIENDIAKATGDIKV 240
Db 181 WASQVDOGANFSEVSNSTFELNSNVTKVSVLVYNTVNTTSCMIENDIAKATGDIKV 240

Qy 241 TESIKRSHLQLNLSKASLCV-SSFFAISWALLPLSPYMLK 282
Db 241 TDSEVKRRSQQLNLSGSPCVFSSAFVAGWALLSLSCCLMLR 283

RESULT 5
Q8K091
ID Q8K091 PRELIMINARY; PRT; 283 AA.
AC Q8K091,
DT 01-OCT-2002 (TrEMBLrel. 22, Created)

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DT 01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
DT 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE Immune costimulatory protein B7-H4.
GN Name=BC032925;
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Uterus;
RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Dege J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.L., Jordan H., Moore T., Max S.I., Wang J., Haieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Udwin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullihy S.J.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Wooley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Kettman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Marra M.A.;
RT "Generation and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences."
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN [2]
RP SEQUENCE FROM N.A.
RC TISSUE=Uterus;
RA Strausberg R.;
RL Submitted (JUN-2002) to the EMBL/GenBank/DBJ databases.
DR EMBL; BC032925; AAH32925.1; -
DR HSP; Q63345; 1PKO.
DR MGD; MGI:3039619; BC032925.
DR InterPro; IPR003599; IG.
DR InterPro; IPR007110; IG-like.
DR SMART; SMO0409; IG; 1.
DR PROSITE; PS50835; IG LIKE; 2.
SQ SEQUENCE 283 AA; 30801 MW; 7E5817417323453B CRC64;

Query Match 87.6%; Score 1253.5; DB 2; Length 283;
Best Local Similarity 87.3%; Pred. No. 3.5e-93;
Matches 247; Conservative 14; Mismatches 21; Indels 1; Gaps 1;

Qy 1 MASLGQLFWSIIIIILAGAIALLIIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP 60
Db 1 MASLGQIIFWSIINIILAGAIALLIIGFGISGRHFTVTTFTSAGNIGEDGILSCTFEP 60

Qy 61 DIKLSDIVIOWLKEGVLGVHEFKEGKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
Db 61 DIKLVGIVOWLKEGIVKGLVHEFKEGKDDLSQQHEMFRGRTAVFADQVIVGNASRLKNV 120

Qy 121 QLTDAGTYKCYIITSKGNANLEYKYGAFSMPVNVYDYNASSETLRCEAPRPFQPTVV 180
Db 121 QLTDAGTYCYIRSSKGNANLEYKYGAFSMPVNVYDYNASSETLRCEAPRPFQPTVA 180

Qy 181 WASQVDOGANFSEVSNSTFELNSNVTKVSVLVYNTVNTTSCMIENDIAKATGDIKV 240
Db 181 WASQVDOGANFSEVSNSTFELNSNVTKVSVLVYNTVNTTSCMIENDIAKATGDIKV 240

Qy 241 TESIKRSHLQLNLSKASLCV-SSFFAISWALLPLSPYMLK 282
Db 241 TDSEVKRRSQQLNLSGSPCVFSSAFVAGWALLSLSCCLMLR 283

RESULT 6
Q6P097

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Db	177	TWVR--DSSGANITNVTIKTFPPALNLT-DVTTELSGAQLDVQYSLIHDTLAQAVGS	233
Qy	239	KVTESIKRSHQLLNKASLCVSSFFAISWALL	273
Qy	60	PDIKLSDIVIQLKEGVGLVHEFEXGKDELSEQDEMFRGRTAVFADQVIVGNASIRLKN	119

Qy 14 SIILLAGAIALIIGFISGRHSITVTTVASAGNIGEDGILSCFEPEDIKLSDIVIOWLK 73
Db 5 SIILLVT---VSLIQSLSERFQVPTKSPVTAIVGSSIELNCHLFPFNAEKMEIRWLR 61
Qy 74 EGVGLGVHEKFGKDELSEQDEMERGRGTAVFADQVIVGNASRLRKNVQLTDAGTYKCYII 133
Db 62 NSFRPYVHLLVINGEDNTRQNEEPRGRTEFLKQNIIGRIGALTIHKVQLSDQGLTYCYFE 121
Qy 134 TSKGKGANLEYKTFGAFSM-PEVNV-DYNASSETLRCEAPRFPQPTVWVASQVDQGANF 191
Db 122 SETNHQAQVELKVAANGLHPPIWVEDYHDKITLNCSSGMPFKPMWQDETGTRES 181
Qy 192 SEVSNTSFELSENENVTMKVSVLYNVTINNTYSQMIENDIAK--ATGDIKVTSEIKR-- 247
Db 182 TEKINVTETGLFHTVSAI-----RIETQAKISCHVRNLDLEDGESSIKFAESFYWRVD 236
Qy 248 RSHLOLINSKASLCVSPFAISWALLPLSP 277
Db 237 RSGISRF-----VLMASCLAVLIVIGTP 261

RESULT 12

Q9BXR1 ID Q9BXR1 PRELIMINARY; PRT; 316 AA.
AC Q9BXR1
DT 01-JUN-2001 (TrEMBLrel. 17, Created)
DT 01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Costimulatory molecule.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=21163383; PubMed=1124528;
RA Chapoval A.I., Ni J., Lau J.S., Wilcox R.A., Flies D.B., Liu D.,
RA Dong H., Sica G.L., Zhu G., Tamada K., Chen L.;
RT "B7-H3: A costimulatory molecule for T cell activation and IFN-gamma
production";
RL Nat. Immun. 2:269-274 (2001).
DR EMBL; AF302102; AAK15438.1; -.
DR HSP; Q63345; IPKO.
DR GO; GO:0008283; P:cell proliferation; NAS.
DR GO; GO:0006955; P:immune response; NAS.
DR InterPro; IPR003599; IG-like.
DR InterPro; IPR007110; IG-like.
DR Pfam; PF00047; IG; 1.
DR SMART; SM00409; IG; 2.
DR PROSITE; PS00835; IG LIKE; 2.
SQ SEQUENCE 316 AA; 33791 MW; PF97007F191CCFA1 CRC64;

Query Match 17.2%; Score 246.5; DB 2; Length 316;
Best Local Similarity 30.2%; Pred. No. 1.2e-11;
Matches 70; Conservative 44; Mismatches 99; Indels 19; Gaps 9;
Qy 21 GAIALIIGFISGRHSITVTTVASAGNIGEDGILSCF--EPDIKLSDIVIOWLKGVLG 78
Db 15 GALLGALWFLUTGALEFVQVPEDPVVALVIGTDATLCCSFSPGSLAQLNLIWQLTQK 74
Qy 79 LVHEKFGKDELSEQDEMERGRGTAVFADQVIVGNASRLRKNVQLTDAGTYKCYIITSKKG 138
Db 75 LVHSFAEQD---QGSAYANRTALFPDLLAQGNASRLRQVRVADSGSTFCF-VSIRDF 129
Qy 139 GNANLEYKTFGAFSMPEVNVVDYN-----ASSETURCEAPRFPQPTVWVASQVDQGANFS 192
Db 130 GSAVSLQVAAPYKSPKSNMTLPKDLRPGDVTITTCSSYRGYPEAEVFW--QDQGVPLT 187
Qy 193 EVSNTSFELSENENVTMKVSVLYNVT-INNTYSQMIENDIAK--ATGDIKVT 241
Db 188 GNVTTIS-QMANEQGLFDVHSLVRVLGANGTYSCLVRNPVLQDAGHSVTIT 238

RESULT 13

Q8NC34 ID Q8NC34 PRELIMINARY; PRT; 388 AA.
AC Q8NC34
DT 01-OCT-2002 (TrEMBLrel. 22, Created)
DT 01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
DT 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE Hypothetical protein FLJ90516.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RA Isogai T., Ota T., Nishikawa T., Hayashi K., Otsuki T., Sugiyama T.,
RA Suzuki Y., Nagai K., Sugano S., Ishii S., Kawai-Hio Y., Saito K.,
RA Yamamoto J., Wakamatsu A., Nakamura Y., Kojima S., Nagahari K.,
RA Masuho Y., Ono T., Okano K., Yoshikawa Y., Aotsuka S., Sasaki N.,
RA Hattori A., Okumura K., Iwayanagi T., Ninomiya K.;
RL Submitted (MAR-2002) to the EMBL/GenBank/DBJ databases.
DR EMBL; AK074997; BAC11344.1; -.
DR HSP; Q63345; IPKO.
DR InterPro; IPR007110; IG-like.
DR InterPro; IPR003598; IG_c2.
DR Pfam; PF00047; IG; 2.
DR SMART; SM00408; IGC2; 1.
DR PROSITE; PS00835; IG LIKE; 3.
SQ SEQUENCE 388 AA; 41768 MW; 44A59B9E3AB3DCD3 CRC64;
Query Match 17.1%; Score 245; DB 2; Length 388;
Best Local Similarity 27.8%; Pred. No. 2e-11;
Matches 72; Conservative 50; Mismatches 101; Indels 36; Gaps 11;
Qy 11 SIISIIILAGAIILIGFGI---SGRHSITVTTVAS-----AGNIGEDGI 53
Db 60 SIIRVLGANGTYSCLVRNPVLQDAGHSVTITPQSPTCGAVEQVPEDPVVALVIGTDAT 119
Qy 54 LSCTF--EPDIKLSDIVIOWLKGVGLVHEKFGKDELSEQDEMERGRGTAVFADQVIVG 111
Db 120 LRCSFSPGSLAQLNLIWQLTDTKLHVSFTGERD---QGSAYANRTALFPDLLAQG 175
Qy 112 NASRLKNVQLTDAGTYKCYIITSKGNANLEYKTFGAFSMPEVNVVDYN-----ASSET 165
Db 176 NASRLQVRVADSGSTFCF-VSIRDFGSAVSLQVAAPYKSPKSNMTLPKDLRPGDVT 234
Qy 166 LRCEAPRFPQPTVWVASQVDQGANFSEVNTSFELSENENVTMKVSVLYNVT-INNTYS 224
Db 235 ITCSYRGYPEAEVFW--QDQGVPLTGNVTTIS-QMANEQGLFDVHSLVRVLGANGTYS 291
Qy 225 CMIENDIAK--ATGDIKVT 241
Db 292 CLVRNPVLQDAGHSVTIT 310

RESULT 14

Q6PSY4 ID Q6PSY4 PRELIMINARY; PRT; 493 AA.
AC Q6PSY4
DT 05-JUL-2004 (TrEMBLrel. 27, Created)
DT 05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
DT 05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
DE Hypothetical protein.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Lung;
RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
 RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
 RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
 RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
 RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
 RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
 RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
 RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
 RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A., Sanchez A.,
 RA Fahey J., Helton E., Kettman M., Madan A.C., Rodrigues S., Sanchez A.,
 RA Whiting J., Madan A.C., Shevchenko Y., Bouffard G.G.,
 RA Brakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
 RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
 RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
 RA Jones S.J., Marra M.A.;
 RT "Generation and initial analysis of more than 15,000 full-length human
 RT and mouse cDNA sequences.";
 RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
 RP [2]
 RP SEQUENCE FROM N.A.
 RC TISSUE=Lung;
 RC Strausberg R.;
 RL Submitted (NOV-2003) to the EMBL/GenBank/DBJ databases.
 DR EMBL; BC062581; AA62581.1; -;
 DR InterPro; IPR003599; IG.
 DR InterPro; IPR007110; IG-like.
 DR InterPro; IPR003597; IG cl.
 DR InterPro; IPR003598; IG c2.
 DR InterPro; IPR003596; IG v.
 DR Pfam; PF00047; IG; 2.
 DR SMART; SM00409; IG; 4.
 DR SMART; SM00407; IGcl; 2.
 DR SMART; SM00408; IGc2; 4.
 DR SMART; SM00406; IGv; 2.
 DR PROSITE; PS50835; IG LIKE; 4.
 KW Hypothetical protein.
 SQ SEQUENCE 493 AA; 52761 MW; CD7C5591CC4822D2 CRC64;

Query Match 17.1%; Score 245; DB 2; Length 493;
 Best Local Similarity 27.8%; Pred. No. 2.8e-11;
 Matches 72; Conservative 50; Mismatches 101; Indels 36; Gaps 11;
 Qy 11 SIISIIILAGAIILIGFI---SGRHSITVTIVAS-----AGNIGEDGI 53
 Db 206 SILRVVLGANGTYSCLVRNPVLQDAHSSVTITPQRSPTGAVEVQVPEDPVVALVGTDAT 265
 Qy 54 LSCTF--EPDIKLSDIVIOWLKEGVLGVHEFKGKDELSEQDEMFRGRTAVFADQVIVG 111
 Db 266 LRCSFSEPGFSLAQLNLWLTDTKQLVHSFTGERD---QGSAYANRTALPDLAQQ 321
 Qy 112 NASLRKNVOLTDAGTVKCYIITSKGNANLEYKTGA-FSMPEVNVDYN-----ASSET 165
 Db 322 NASLRQVRVADEGSFTCF-VSIRDFGSAVSLQVAAPYKPSMTLEPNKDLRPGDTVT 380
 Qy 166 LRCEAPRWFPPQPTVVWASQVDQGANFSEVNTSFEINSENVTMKVSVLYNVT-INNTYS 224
 Db 381 ITCSSYRGYPEAEVFW--QDQGGVPLTGNVTTTS-QMANEQGLFDVHSLRVVLGANGTYS 437
 Qy 225 CMIENDIAK--ATGDIKVT 241
 Db 438 CLVRNPVLQDAGHSVTIT 456

RESULT 15

Q8NCB6 PRELIMINARY; PRT; 533 AA.
 AC Q8NCB6
 DT 01-OCT-2002 (TrEMBLrel. 22, Created)
 DT 01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
 DT 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
 DE Hypothetical protein FLJ90368.
 OS Homo sapiens (Human)
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

OC Mammalia; Euthera; Primates; Catarrhini; Homnidae; Homo.
 OX NCBI_TaxID=9606;
 RN [1]
 RP SEQUENCE FROM N.A.
 RA Isegai T., Ota T., Nishikawa T., Hayashi K., Otsuki T., Sugiyama T.,
 RA Suzuki Y., Nagai K., Sugano S., Ishii S., Kawai-Hio Y., Saito K.,
 RA Yamamoto J., Wakamatsu A., Nakamura Y., Kojima S., Nagahari K.,
 RA Masuho Y., Ono T., Okano K., Yoshikawa Y., Aotsuka S., Sasaki N.,
 RA Hattori A., Okumura K., Iwayanagi T., Ninomiya K.;
 RL Submitted (MAR-2002) to the EMBL/GenBank/DBJ databases.
 DR EMBL; AK074849; BAC11243.1; -;
 DR HSP; Q63345; 1PKO.
 DR InterPro; IPR003599; IG.
 DR InterPro; IPR007110; IG-like.
 DR Pfam; PF00047; IG; 2.
 DR SMART; SM00409; IG; 4.
 DR PROSITE; PS50835; IG LIKE; 4.
 SQ SEQUENCE 533 AA; 57179 MW; FC7E3E3A84F56A56 CRC64;
 Query Match 17.1%; Score 245; DB 2; Length 533;
 Best Local Similarity 27.8%; Pred. No. 3.1e-11;
 Matches 72; Conservative 50; Mismatches 101; Indels 36; Gaps 11;
 Qy 11 SIISIIILAGAIILIGFI---SGRHSITVTIVAS-----AGNIGEDGI 53
 Db 206 SILRVVLGANGTYSCLVRNPVLQDAHSSVTITPQRSPTGAVEVQVPEDPVVALVGTDAT 265
 Qy 54 LSCTF--EPDIKLSDIVIOWLKEGVLGVHEFKGKDELSEQDEMFRGRTAVFADQVIVG 111
 Db 266 LRCSFSEPGFSLAQLNLWLTDTKQLVHSFTGERD---QGSAYANRTALPDLAQQ 321
 Qy 112 NASLRKNVOLTDAGTVKCYIITSKGNANLEYKTGA-FSMPEVNVDYN-----ASSET 165
 Db 322 NASLRQVRVADEGSFTCF-VSIRDFGSAVSLQVAAPYKPSMTLEPNKDLRPGDTVT 380
 Qy 166 LRCEAPRWFPPQPTVVWASQVDQGANFSEVNTSFEINSENVTMKVSVLYNVT-INNTYS 224
 Db 381 ITCSSYRGYPEAEVFW--QDQGGVPLTGNVTTTS-QMANEQGLFDVHSLRVVLGANGTYS 437
 Qy 225 CMIENDIAK--ATGDIKVT 241
 Db 438 CLVRNPVLQDAGHSVTIT 456
 Search completed: April 19, 2005, 07:15:01
 Job time : 93 secs

Result No.	Query %			DB	ID	Description
	Score	Match	Length			
1	1431	100.0	282	9	US-09-778-320-208	Sequence 208, App
2	1431	100.0	282	9	US-09-850-178-33	Sequence 33, Appl
3	1431	100.0	282	9	US-09-877-065-8	Sequence 8, Appl
4	1431	100.0	282	9	US-09-989-723-291	Sequence 291, App
5	1431	100.0	282	9	US-09-989-723-291	Sequence 291, App
6	1431	100.0	282	9	US-09-989-273-291	Sequence 291, App
7	1431	100.0	282	9	US-09-989-727-291	Sequence 291, App
8	1431	100.0	282	9	US-09-910-689-208	Sequence 208, App
9	1431	100.0	282	9	US-09-989-731-291	Sequence 291, App
10	1431	100.0	282	9	US-09-884-441-393	Sequence 393, App
11	1431	100.0	282	9	US-09-989-733-291	Sequence 291, App
12	1431	100.0	282	9	US-09-991-073-291	Sequence 291, App
13	1431	100.0	282	9	US-09-990-441-291	Sequence 291, App

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121 QLTDAQYKCYIITSKGNANLEYKTFGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180
121 QLTDAQYKCYIITSKGNANLEYKTFGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180
181 WASQVQGANFSEVNTSFEINSENVTMKVSVLYNNTTSCMIENDIAKATGDIKV 240
181 WASQVQGANFSEVNTSFEINSENVTMKVSVLYNNTTSCMIENDIAKATGDIKV 240
241 TESEIKRRSHLQLLNSKASLCVSSFFAISWALLPLSPYMLK 282
241 TESEIKRRSHLQLLNSKASLCVSSFFAISWALLPLSPYMLK 282

RESULT 2
US-09-850-178-33
; Sequence 33, Application US/09850178
; Patent No. US20020034749A1
; GENERAL INFORMATION:
; APPLICANT: Abbott Laboratories
; APPLICANT: Billing-Medel, Patricia A.
; APPLICANT: Cohen, Maurice
; APPLICANT: Colipitts, Tracey L.
; APPLICANT: Friedman, Paula N.
; APPLICANT: Russell, John C.
; APPLICANT: Granados, Edward N.
; APPLICANT: Hodges, Steven C.
; APPLICANT: Klass, Michael R.
; APPLICANT: Kratochvil, Jon D.
; APPLICANT: Roberts-Rapp, Lisa
; APPLICANT: Stroupe, Stephen D.
; APPLICANT: Gordon, Juliana
; TITLE OF INVENTION: REAGENTS AND METHODS USEFUL FOR
; DETECTING DISEASES OF THE BREAST
; FILE REFERENCE: 6251.US.P1
; CURRENT APPLICATION NUMBER: US/09/850,178
; PRIOR FILING DATE: 2001-05-07
; PRIOR APPLICATION NUMBER: US 08/972,376
; PRIOR FILING DATE: 1997-11-18
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 33
; LENGTH: 282
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-850-178-33

Query Match 100.0%; Score 1431; DB 9; Length 282;
Best Local Similarity 100.0%; Pred. No. 2.7e-112;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MASLGOILFWSIISIIILAGAILIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60
DB 1 MASLGOILFWSIISIIILAGAILIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60
QY 61 DIKLSDIVIOWLKEGVLGVHFEKGEKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
DB 61 DIKLSDIVIOWLKEGVLGVHFEKGEKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
QY 121 QLTDAQYKCYIITSKGNANLEYKTFGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180
DB 121 QLTDAQYKCYIITSKGNANLEYKTFGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180
QY 181 WASQVQGANFSEVNTSFEINSENVTMKVSVLYNNTTSCMIENDIAKATGDIKV 240
DB 181 WASQVQGANFSEVNTSFEINSENVTMKVSVLYNNTTSCMIENDIAKATGDIKV 240
QY 241 TESEIKRRSHLQLLNSKASLCVSSFFAISWALLPLSPYMLK 282
DB 241 TESEIKRRSHLQLLNSKASLCVSSFFAISWALLPLSPYMLK 282

RESULT 3
US-09-850-178-33
; Sequence 33, Application US/09850178
; Patent No. US20020034749A1
; GENERAL INFORMATION:
; APPLICANT: Abbott Laboratories
; APPLICANT: Billing-Medel, Patricia A.
; APPLICANT: Cohen, Maurice
; APPLICANT: Colipitts, Tracey L.
; APPLICANT: Friedman, Paula N.
; APPLICANT: Russell, John C.
; APPLICANT: Granados, Edward N.
; APPLICANT: Hodges, Steven C.
; APPLICANT: Klass, Michael R.
; APPLICANT: Kratochvil, Jon D.
; APPLICANT: Roberts-Rapp, Lisa
; APPLICANT: Stroupe, Stephen D.
; APPLICANT: Gordon, Juliana
; TITLE OF INVENTION: REAGENTS AND METHODS USEFUL FOR
; DETECTING DISEASES OF THE BREAST
; FILE REFERENCE: 6251.US.P1
; CURRENT APPLICATION NUMBER: US/09/850,178
; PRIOR FILING DATE: 2001-05-07
; PRIOR APPLICATION NUMBER: US 08/972,376
; PRIOR FILING DATE: 1997-11-18
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 33
; LENGTH: 282
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-850-178-33

Query Match 100.0%; Score 1431; DB 9; Length 282;
Best Local Similarity 100.0%; Pred. No. 2.7e-112;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MASLGOILFWSIISIIILAGAILIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60
DB 1 MASLGOILFWSIISIIILAGAILIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60
QY 61 DIKLSDIVIOWLKEGVLGVHFEKGEKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
DB 61 DIKLSDIVIOWLKEGVLGVHFEKGEKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
QY 121 QLTDAQYKCYIITSKGNANLEYKTFGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180
DB 121 QLTDAQYKCYIITSKGNANLEYKTFGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180
QY 181 WASQVQGANFSEVNTSFEINSENVTMKVSVLYNNTTSCMIENDIAKATGDIKV 240
DB 181 WASQVQGANFSEVNTSFEINSENVTMKVSVLYNNTTSCMIENDIAKATGDIKV 240
QY 241 TESEIKRRSHLQLLNSKASLCVSSFFAISWALLPLSPYMLK 282
DB 241 TESEIKRRSHLQLLNSKASLCVSSFFAISWALLPLSPYMLK 282
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US-09-877-065-8
; Sequence 8, Application US/09877065
; Patent No. US20020051990A1
; GENERAL INFORMATION:
; APPLICANT: OPLE, ERIC
; APPLICANT: MCLACHLAN, KAREN
; APPLICANT: HEARD, CHERYL J.
; TITLE OF INVENTION: NOVEL GENE TARGETS AND LIGANDS THAT BIND THERETO FOR
; TREATMENT AND DIAGNOSIS OF OVARIAN CARCINOMAS
; FILE REFERENCE: 037003-0280631
; CURRENT APPLICATION NUMBER: US/09/877,065
; CURRENT FILING DATE: 2001-06-11
; PRIOR APPLICATION NUMBER: 60/210,451
; PRIOR FILING DATE: 2000-06-09
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 8
; LENGTH: 282
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-877-065-8

Query Match 100.0%; Score 1431; DB 9; Length 282;
Best Local Similarity 100.0%; Pred. No. 2.7e-112;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MASLGOILFWSIISIIILAGAILIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60
DB 1 MASLGOILFWSIISIIILAGAILIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60
QY 61 DIKLSDIVIOWLKEGVLGVHFEKGEKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
DB 61 DIKLSDIVIOWLKEGVLGVHFEKGEKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
QY 121 QLTDAQYKCYIITSKGNANLEYKTFGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180
DB 121 QLTDAQYKCYIITSKGNANLEYKTFGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180
QY 181 WASQVQGANFSEVNTSFEINSENVTMKVSVLYNNTTSCMIENDIAKATGDIKV 240
DB 181 WASQVQGANFSEVNTSFEINSENVTMKVSVLYNNTTSCMIENDIAKATGDIKV 240
QY 241 TESEIKRRSHLQLLNSKASLCVSSFFAISWALLPLSPYMLK 282
DB 241 TESEIKRRSHLQLLNSKASLCVSSFFAISWALLPLSPYMLK 282

RESULT 4
US-09-989-722-291
; Sequence 291, Application US/09989722
; Patent No. US20020072067A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Nepier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
US-09-989-722-291
```

APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William I.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE REFERENCE: P2730P1C63
CURRENT APPLICATION NUMBER: US/09/989,722
CURRENT FILING DATE: 2001-11-19
PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/065186
PRIOR FILING DATE: 1997-11-12
PRIOR APPLICATION NUMBER: 60/065311
PRIOR FILING DATE: 1997-11-13
PRIOR APPLICATION NUMBER: 60/066770
PRIOR FILING DATE: 1997-11-24
PRIOR APPLICATION NUMBER: 60/075945
PRIOR FILING DATE: 1998-02-25
PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/083322
PRIOR FILING DATE: 1998-04-28
PRIOR APPLICATION NUMBER: 60/084600
PRIOR FILING DATE: 1998-05-07
PRIOR APPLICATION NUMBER: 60/087106
PRIOR FILING DATE: 1998-05-28
PRIOR APPLICATION NUMBER: 60/087607
PRIOR FILING DATE: 1998-06-02
PRIOR APPLICATION NUMBER: 60/087609
PRIOR FILING DATE: 1998-06-02
PRIOR APPLICATION NUMBER: 60/087759
PRIOR FILING DATE: 1998-06-02
PRIOR APPLICATION NUMBER: 60/087827
PRIOR FILING DATE: 1998-06-03
PRIOR APPLICATION NUMBER: 60/088021
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Query Match 100.0%; Score 1431; DB 9; Length 282;

Best Local Similarity 100.0%; Pred. No. 2.7e-112; Mismatches 0; Indels 0; Gaps 0;
Matches 282; Conservative 0;

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Qy	241	TESEIKRSHLOLNSKASLCVSSFFAISWALLPLSPYLMK	282
Db	241	TESEIKRSHLOLNSKASLCVSSFFAISWALLPLSPYLMK	282

RESULT 5

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; Sequence 291, Application US/09989723
; Patent No. US20020072092A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Deanoysers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
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; APPLICANT: Grimaldi, J. Christopher

;; APPLICANT: Gurney, Austin L.
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;; APPLICANT: Tumas, Daniel
;; APPLICANT: Watanabe, Colin K.
;; APPLICANT: Williams, P. Mickey
;; APPLICANT: Wood, William I.
;; APPLICANT: Zhang, Zemin
;; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
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Qy 181 WASQVDOGANFSEVSNSTSFELNSENVTMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
Db 181 WASQVDOGANFSEVSNSTSFELNSENVTMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
Qy 241 TESEIKRSHQLNLNSKASLCVSSFFAISWALLPLSPYLMK 282
Db 241 TESEIKRSHQLNLNSKASLCVSSFFAISWALLPLSPYLMK 282

RESULT 6

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; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David

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;; PRIOR FILING DATE: 1998-07-07
;; PRIOR APPLICATION NUMBER: 60/092182
;; PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 1431; DB 9; Length 282;
Best Local Similarity 100.0%; Pred. No. 2,7e-112;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASIGQILFWSIISIIILLAGAIALIIIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP 60
Db 1 MASIGQILFWSIISIIILLAGAIALIIIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP 60

Qy 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
Db 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120

Qy 121 QLTDAGTYKCVIITTSKGNANLEYKTKGAFSMPENVVNDYNASSETLRCEAPRFPQPTVV 180
Db 121 QLTDAGTYKCVIITTSKGNANLEYKTKGAFSMPENVVNDYNASSETLRCEAPRFPQPTVV 180

Qy 181 WASQVDOGANFSEVSNSTSFELNSENVTKVSVLYNVYNTINNTYSCMIENDIAKATGDIKV 240
Db 181 WASQVDOGANFSEVSNSTSFELNSENVTKVSVLYNVYNTINNTYSCMIENDIAKATGDIKV 240

Qy 241 TSEIKRSHQLLNKASLCVSSFFAISWALLPLSPVLMK 282
Db 241 TSEIKRSHQLLNKASLCVSSFFAISWALLPLSPVLMK 282

RESULT 7

US-09-989-727-291
; Sequence 291, Application US/09989727
; Patent No. US20020072497A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Grittisen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kijavini, Ivar J.
; APPLICANT: Napier, Mary A.
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; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: P2730P1C65
; CURRENT APPLICATION NUMBER: US/09/989,727
; CURRENT FILING DATE: 2001-11-19
; PRIOR APPLICATION NUMBER: 60/049787
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; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 1431; DB 9; Length 282;

Best Local Similarity 100.0%; Pred. No. 2.7e-112;

Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASIGQILFWSIISIIILAGALALIIIGFISGRHSITVTTVASAGNIGDGLSCTFEP 60

Db 1 MASIGQILFWSIISIIILAGALALIIIGFISGRHSITVTTVASAGNIGDGLSCTFEP 60

Qy 61 DIKLSDIVIQWLKEGVLGVHFEKKGKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120

Db 61 DIKLSDIVIQWLKEGVLGVHFEKKGKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120

Qy 121 QLTDACTYKCYIITSKGKNANLEYKTGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180

Db 121 QLTDACTYKCYIITSKGKNANLEYKTGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180

Qy	181	WASQVDQAGNFSEVNTSFELNSENVTWKVSVLYNVTNNYTSY	240
Db	181	WASQVDQAGNFSEVNTSFELNSENVTWKVSVLYNVTNNYTSY	240
Qy	241	TESBIKRSHIQLNLSKSLCVSSFFAISWALLPLSPYIMLK	282
Db	241	TESBIKRSHIQLNLSKSLCVSSFFAISWALLPLSPYIMLK	282

RESULT 8

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US-09-910-689-208
; Sequence 208, Application US/09910689
; Patent No. US20020081609A1
; GENERAL INFORMATION:
; APPLICANT: Dillon, Davin C.
; APPLICANT: Day, Craig H.
; APPLICANT: Jiang, Yuqiu
; APPLICANT: Houghton, Raymond L.
; APPLICANT: Mitcham, Jennifer
; APPLICANT: Wang, Tongtong
; APPLICANT: McNeill, Patricia D.
; APPLICANT: Harlocker, Susan L.
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
; TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
; FILE REFERENCE: 210121.491C6
; CURRENT APPLICATION NUMBER: US/09/910,689
; CURRENT FILING DATE: 2001-07-20
; NUMBER OF SEQ ID NOS: 307
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 208
; LENGTH: 282
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-910-689-208

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Query Match      100.0%; Score 1431; DB 9; Length 282;
Best Local Similarity 100.0%; Pred. No. 2.7e-112;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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Db	61	DIKLSDIVIOWLKEGVGLV	HEFKEGKDELSQDEMFGRGTAFADQVTVGNASLRLKNV	120
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Db	121	QLTDAGTYKCVIIITSK	GKGNANLEYKTAFSMPENVVDYNASSETLRCPEARWFPQPTVV	180
Qy	181	WASQVDOGANFSEVNS	TSFELSENVTWKVSVLVNVTINNTYSCHIENDIAKATGDIKV	240
Db	181	WASQVDOGANFSEVNS	TSFELSENVTWKVSVLVNVTINNTYSCHIENDIAKATGDIKV	240
Qy	241	TESIIKRSHIOLANSKAS	LCVSFFAISWALLPLSPYMLK	282
Db	241	TESIIKRSHIOLANSKAS	LCVSFFAISWALLPLSPYMLK	282

RESULT 9

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US-09-989-731-291
; Sequence 291, Application US/09989731
; Patent No. US20020103125A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Deenoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman

```

1	APPLICANT:	Gerber, Hanspeter	
2	APPLICANT:	Gerritsen, Mary E.	
3	APPLICANT:	Goddard, Audrey	
4	APPLICANT:	Godowski, Paul J.	
5	APPLICANT:	Grimaldi, J. Christopher	
6	APPLICANT:	Gurney, Austin L.	
7	APPLICANT:	Kljasin, Ivar J.	
8	APPLICANT:	Napier, Mary A.	
9	APPLICANT:	Pan, James	
10	APPLICANT:	Paoni, Nicholas F.	
11	APPLICANT:	Roy, Margaret Ann	
12	APPLICANT:	Stewart, Timothy A.	
13	APPLICANT:	Tumas, Daniel	
14	APPLICANT:	Watanabe, Colin K.	
15	APPLICANT:	Williams, P. Mickey	
16	APPLICANT:	Wood, William I.	
17	APPLICANT:	Zhang, Zemin	
18	TITLE OF INVENTION:	Secreted and Transmembrane Polypeptides and Nucleic	
19	TITLE OF INVENTION:	Acids Encoding the Same	
20	FILE REFERENCE:	P2730PIC70	
21	CURRENT APPLICATION NUMBER:	US/09/989,731	
22	CURRENT FILING DATE:	2001-11-20	
23	PRIOR APPLICATION NUMBER:	60/049787	
24	PRIOR FILING DATE:	1997-06-16	
25	PRIOR APPLICATION NUMBER:	60/062250	
26	PRIOR FILING DATE:	1997-10-17	
27	PRIOR APPLICATION NUMBER:	60/065186	
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29	PRIOR APPLICATION NUMBER:	60/065311	
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34	PRIOR FILING DATE:	1998-02-25	
35	PRIOR APPLICATION NUMBER:	60/078910	
36	PRIOR FILING DATE:	1998-03-20	
37	PRIOR APPLICATION NUMBER:	60/083322	
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; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 1431; DB 9; Length 282;
Best Local Similarity 100.0%; Pred. No. 2.7e-112;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy	61	DIKLSDIVIQWLKEGVLGVHFEKGEKDELSEODEMFRGRTAVFADQVIVGNASRLKNV	120
Db	61	DIKLSDIVIQWLKEGVLGVHFEKGEKDELSEODEMFRGRTAVFADQVIVGNASRLKNV	120
Qy	121	QLTDAITYKCYIITSKGKNANLEYKTGAFSPMEVNVVDYNASSETLRCEAPRFPPTVV	180
Db	121	QLTDAITYKCYIITSKGKNANLEYKTGAFSPMEVNVVDYNASSETLRCEAPRFPPTVV	180
Qy	181	WASQVDQGANFSEVSNSTSFELNSENVTMKVSVLYNVNTINNTYSCTMIENDIAKATGDIKV	240
Db	181	WASQVDQGANFSEVSNSTSFELNSENVTMKVSVLYNVNTINNTYSCTMIENDIAKATGDIKV	240
Qy	241	TESEIKRRSHLQLNLSKASLCVSSFFAISWALLPLSPYLMK	282
Db	241	TESEIKRRSHLQLNLSKASLCVSSFFAISWALLPLSPYLMK	282

RESULT 10
US-09-884-441-393
; Sequence 393, Application US/09884441

```

; Patent No. US20020119158A1
; GENERAL INFORMATION:
; APPLICANT: Algate, Paul A.
; APPLICANT: Carter, Darrick
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
; TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
; FILE REFERENCE: 210121.462C7
; CURRENT APPLICATION NUMBER: US/09/884,441
; CURRENT FILING DATE: 2001-06-18
; NUMBER OF SEQ ID NOS: 489
; SOFTWARE: FastSeq for Windows Version 3.0
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; LENGTH: 282
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-884-441-393

Query Match      100.0%; Score 1431; DB 9; Length 282;
Best Local Similarity 100.0%; Pred. No. 2.7e-112;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db      61 DIKLSDIVIOWKEGVLGLVHEFKEGKDELSEQDEMFRGRTAVFADQVIVGNASLRKNV 120

Qy      121 QLTAGTYKCVIITSKGKGNANLEYKTCGAFSMPEVNVVDYNASSETLRCEAPRWFPTTV 180
Db      121 QLTAGTYKCVIITSKGKGNANLEYKTCGAFSMPEVNVVDYNASSETLRCEAPRWFPTTV 180

Qy      181 WASQVDQANFSEVNSFTSELSNVTKVSVLYVNTTNTYSCMIENDIAKATGDIKV 240
Db      181 WASQVDQANFSEVNSFTSELSNVTKVSVLYVNTTNTYSCMIENDIAKATGDIKV 240

Qy      241 TESEIKRRSHQLLNKASCLVSSFFAISWALLPLSPYMLK 282
Db      241 TESEIKRRSHQLLNKASCLVSSFFAISWALLPLSPYMLK 282

RESULT 11
US-09-989-732-291
; Sequence 291, Application US/09989732
; Patent No. US20020123463A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Napier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: P2730P1C57
; CURRENT APPLICATION NUMBER: US/09/989,732
; CURRENT FILING DATE: 2001-11-19
; PRIOR APPLICATION NUMBER: 60/049787
; PRIOR FILING DATE: 1997-06-16
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; PRIOR APPLICATION NUMBER: 60/091978
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 1431; DB 9; Length 282;

Best Local Similarity 100.0%; Pred. No. 2.7e-112; Mismatches 0; Indels 0; Gaps 0;
Matches 282; Conservative 0

Qy	1	MASLGQILFWSIISIIILAGATAIILGFGISGRHSITVTTVASAGNIGEDGILSCTFEP	60
Db	1	MASLGQILFWSIISIIILAGATAIILGFGISGRHSITVTTVASAGNIGEDGILSCTFEP	60
Qy	61	DIKLSDIVIOWLKEGVLGVHFEKKGDELSEQDEMFRGRTAVFADQVIVGNASRLKNV	120
Db	61	DIKLSDIVIOWLKEGVLGVHFEKKGDELSEQDEMFRGRTAVFADQVIVGNASRLKNV	120
Qy	121	QLTDAGTYKCYIITSKGGNANLEYKTGAFSMPEVNVVDYNASSETLCEAPRWPQPTVV	180
Db	121	QLTDAGTYKCYIITSKGGNANLEYKTGAFSMPEVNVVDYNASSETLCEAPRWPQPTVV	180
Qy	181	WASQVDOGANFSEVSNTSFELNSENVTMKVSVLYNVNTINNTYSCTMIENDIAKATGDIKV	240
Db	181	WASQVDOGANFSEVSNTSFELNSENVTMKVSVLYNVNTINNTYSCTMIENDIAKATGDIKV	240
Qy	241	TESEIKRRSHLQLLNKSKASLCVSSFFAISWALLPLSPYMLK	282
Db	241	TESEIKRRSHLQLLNKSKASLCVSSFFAISWALLPLSPYMLK	282

RESULT 12

US-09-991-073-291
; Sequence 291, Application US/09991073
; Patent No. US2002012756A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kijavini, Ivar J.
; APPLICANT: Napier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.

APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Watanabe, Colin K.
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William I.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C15
CURRENT APPLICATION NUMBER: US/09/991,073
CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-10-17
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PRIOR FILING DATE: 1997-11-12
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PRIOR APPLICATION NUMBER: 60/075945
PRIOR FILING DATE: 1998-02-25
PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1998-03-20
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PRIOR FILING DATE: 1998-04-28
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PRIOR APPLICATION NUMBER: 60/087106
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Query Match 100.0%; Score 1431; DB 9; Length 282;
Best Local Similarity 100.0%; Pred. No. 2.7e-112;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASLGQLFWSIIIIILAGAIALLIIFGIGSRHSITVTTVASAGNIGEDGILSCTFEP 60
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Qy 121 QLTDAGTYKCYIITSKGNANLEYKTGAFSMPENVVDYNASSETLRCEAPRFPPTVV 180
Db 121 QLTDAGTYKCYIITSKGNANLEYKTGAFSMPENVVDYNASSETLRCEAPRFPPTVV 180

Qy 181 WASQVDOGAFSEVSNSTFELNSENVTKVSVLYNVTINNTYSCMIENDIAKATGDIKV 240
Db 181 WASQVDOGAFSEVSNSTFELNSENVTKVSVLYNVTINNTYSCMIENDIAKATGDIKV 240

Qy 241 TSEIKRSHQLLNSKASLCVSSFALSPVLMK 282
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RESULT 13
US-09-990-442-291
; Sequence 291, Application US/09990442
; Patent No. US20020132252A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter

; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
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; APPLICANT: Paoni, Nicholas F.
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; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: P2730P1C8
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; CURRENT FILING DATE: 2001-11-14
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;	PRIOR FILING DATE:	1998-07-09	

Query Match 100.0%; Score 1431; DB 9; Length 282;
Best Local Similarity 100.0%; Pred. No. 2.7e-112;
Matches 282; Conservative 0; Mismatches 0; Indels 0

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121	Qy	QLTDAGTYKCYIIITSGKGKNANLEYKTGAFSPMEPVNVVDYNASSTLRCEAPRPFPQPVV	180
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181	Qy	WASQVDOGANFSEVSNITSFELNSENVTMKVSVLYNYVTINNTYSCTMIENDIAKATGDIKV	240
181	Db	WASQVDOGANFSEVSNITSFELNSENVTMKVSVLYNYVTINNTYSCTMIENDIAKATGDIKV	240
241	Qy	TESEIKRRSHLOLLNKSASLCVCSFFAISWALLPLSPYLMK	282
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RESULT 14

RESULT 14
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; Sequence 291, Application US/09991163
; Patent No. US20020132253A1

GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
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; APPLICANT: Gerritsen, Mary E.
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; APPLICANT: Gurney, Austin L.
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; APPLICANT: Napier, Mary A.
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; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
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: APPLICANT: Williams, P. Mickey
: APPLICANT: Wood, William I.
: APPLICANT: Zhang, Zemin
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1 PRIOR FILING DATE: 1998-06-24
1 PRIOR APPLICATION NUMBER: 60/090445
1 PRIOR FILING DATE: 1998-06-24
1 PRIOR APPLICATION NUMBER: 60/090472
1 PRIOR FILING DATE: 1998-06-24
1 PRIOR APPLICATION NUMBER: 60/090535
1 PRIOR FILING DATE: 1998-06-24
1 PRIOR APPLICATION NUMBER: 60/090540
1 PRIOR FILING DATE: 1998-06-24
1 PRIOR APPLICATION NUMBER: 60/090542
1 PRIOR FILING DATE: 1998-06-24
1 PRIOR APPLICATION NUMBER: 60/090557
1 PRIOR FILING DATE: 1998-06-24
1 PRIOR APPLICATION NUMBER: 60/090676
1 PRIOR FILING DATE: 1998-06-25
1 PRIOR APPLICATION NUMBER: 60/090678
1 PRIOR FILING DATE: 1998-06-25
1 PRIOR APPLICATION NUMBER: 60/090690
1 PRIOR FILING DATE: 1998-06-25
1 PRIOR APPLICATION NUMBER: 60/090694
1 PRIOR FILING DATE: 1998-06-25
1 PRIOR APPLICATION NUMBER: 60/090695
1 PRIOR FILING DATE: 1998-06-25
1 PRIOR APPLICATION NUMBER: 60/090696
1 PRIOR FILING DATE: 1998-06-25
1 PRIOR APPLICATION NUMBER: 60/090862
1 PRIOR FILING DATE: 1998-06-26
1 PRIOR APPLICATION NUMBER: 60/090863
1 PRIOR FILING DATE: 1998-06-26
1 PRIOR APPLICATION NUMBER: 60/091360
1 PRIOR FILING DATE: 1998-07-01
1 PRIOR APPLICATION NUMBER: 60/091478
1 PRIOR FILING DATE: 1998-07-02
1 PRIOR APPLICATION NUMBER: 60/091544
1 PRIOR FILING DATE: 1998-07-01
1 PRIOR APPLICATION NUMBER: 60/091519
1 PRIOR FILING DATE: 1998-07-02
1 PRIOR APPLICATION NUMBER: 60/091626
1 PRIOR FILING DATE: 1998-07-02
1 PRIOR APPLICATION NUMBER: 60/091633
1 PRIOR FILING DATE: 1998-07-02
1 PRIOR APPLICATION NUMBER: 60/091778
1 PRIOR FILING DATE: 1998-07-07
1 PRIOR APPLICATION NUMBER: 60/091982
1 PRIOR FILING DATE: 1998-07-07
1 PRIOR APPLICATION NUMBER: 60/092182
1 PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 1431; DB 9; Length 282;

Best Local Similarity 100.0%; Pred. No. 2.7e-112;

Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASLGQLFWSIISIILLAGAILIIFGIGSRHSITVTTVASAGNIGEDGILSCTFEP 60
|||||

Db 1 MASLGQLFWSIISIILLAGAILIIFGIGSRHSITVTTVASAGNIGEDGILSCTFEP 60
|||||

Qy 61 DIKLSDIVIQWLKEGVLGVHEFKEGKDELSEODEMPRGRTAVFADQVIVGNASRLKNV 120
|||||

Db 61 DIKLSDIVIQWLKEGVLGVHEFKEGKDELSEODEMPRGRTAVFADQVIVGNASRLKNV 120
|||||

Qy	121	QLTDAGTYKCYII	TSKGGKGNANLEYK	GAFSMPENVNDYNASSETLR	CEAPRWPPTVV	180
Db	121	QLTDAGTYKCYII	TSKGGKGNANLEYK	GAFSMPENVNDYNASSETLR	CEAPRWPPTVV	180
Qy	181	WASOVDOGANFSEV	SNTSFELNSENVTKVSV	LVNVTINNNTYSCMIENDI	AKATGDIKV	240
Db	181	WASOVDOGANFSEV	SNTSFELNSENVTKVSV	LVNVTINNNTYSCMIENDI	AKATGDIKV	240
Qy	241	TESEIKRRSHLQL	LNKASLCVSSFFAISW	ALLPLSPYMLK		282
Db	241	TESEIKRRSHLQL	LNKASLCVSSFFAISW	ALLPLSPYMLK		282

Search completed: April 19, 2005, 07:30:09
Job time : 492 secs

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GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: April 19, 2005, 01:25:19 ; Search time 90 seconds
(without alignments)
1211.849 Million cell updates/sec

Title: US-10-773-715-6

Perfect score: 1431

Sequence: 1 MASLGQLFWSIIIIIIILA.....SSPFAISWALLPLSPYMLK 282

Scoring table:

BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 2105692 seqs, 386760381 residues

Total number of hits satisfying chosen parameters: 2105692

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

A_Geneseq_16Dec04:*

- 1: Geneseqp1980s:*
- 2: Geneseqp1990s:*
- 3: Geneseqp2000s:*
- 4: Geneseqp2001s:*
- 5: Geneseqp2002s:*
- 6: Geneseqp2003as:*
- 7: Geneseqp2003bs:*
- 8: Geneseqp2004s:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	1431	100.0	282	3	AY66719 Membrane-
2	1431	100.0	282	3	AB12557 Human ova
3	1431	100.0	282	4	AU29132 Human PRO
4	1431	100.0	282	4	AB87555 Human PRO
5	1431	100.0	282	4	AB99204 Human ova
6	1431	100.0	282	4	AB65242 Human PRO
7	1431	100.0	282	5	AE20311 Human B7-
8	1431	100.0	282	5	ABG96445 Human ova
9	1431	100.0	282	5	AU77766 Tumour as
10	1431	100.0	282	5	ABG95880 Human sec
11	1431	100.0	282	5	AU76536 Tumour-as
12	1431	100.0	282	5	ABP30901 OBE prote
13	1431	100.0	282	5	ABP76274 Breast BS
14	1431	100.0	282	5	AE18336 Human B7-
15	1431	100.0	282	5	ABB09879 Amino aci
16	1431	100.0	282	5	AE19013 Human B7-
17	1431	100.0	282	6	ABU58508 Human PRO
18	1431	100.0	282	6	ABU88056 Novel hum
19	1431	100.0	282	6	ABU84371 Human sec
20	1431	100.0	282	6	ABR66245 Human sec
21	1431	100.0	282	6	ABR65635 Human sec
22	1431	100.0	282	6	ABU99575 Human sec
23	1431	100.0	282	6	ABU58057 Human PRO
24	1431	100.0	282	6	ABU59135 Novel hum
25	1431	100.0	282	6	ABU82647 Human sec

26	1431	100.0	282	6	ABU82814	Human PRO
27	1431	100.0	282	6	ABU89935	Novel hum
28	1431	100.0	282	6	ABR68184	Human sec
29	1431	100.0	282	6	ABU60566	Human sec
30	1431	100.0	282	6	ABU96237	Novel hum
31	1431	100.0	282	6	ABU92668	Human sec
32	1431	100.0	282	6	ABO08745	Human sec
33	1431	100.0	282	6	ABO02797	Human sec
34	1431	100.0	282	6	ABR74951	Human sec
35	1431	100.0	282	6	ABR94713	Human sec
36	1431	100.0	282	6	ABU13948	Human PRO
37	1431	100.0	282	6	ABU85686	Human PRO
38	1431	100.0	282	6	ABU98846	Novel hum
39	1431	100.0	282	6	ABU98061	Novel hum
40	1431	100.0	282	6	ABU91767	Novel hum
41	1431	100.0	282	6	ABU89460	Human PRO
42	1431	100.0	282	6	ABU86301	Human sec
43	1431	100.0	282	6	ABU67514	Human sec
44	1431	100.0	282	6	ABU80542	Human PRO
45	1431	100.0	282	6	ABU72533	Novel hum

ALIGNMENTS

RESULT 1

AY66719	AY66719 standard; protein; 282 AA.
XX	XX
AC	AY66719;
XX	XX
DT	05-APR-2000 (first entry)
XX	XX
DE	Membrane-bound protein PRO1291.
XX	XX
KW	Membrane-bound polypeptide; PRO polypeptide; LDL receptor; TIE ligand;
KW	pharmaceutical; receptor immunoadhesin; gene mapping.
XX	XX
OS	Homo sapiens.
XX	XX
PN	WO9963088-A2.
XX	XX
PD	09-DEC-1999.
XX	XX
PF	02-JUN-1999; 99WO-US012252.
XX	XX
PR	02-JUN-1998; 98US-0087607P.
PR	02-JUN-1998; 98US-0087609P.
PR	02-JUN-1998; 98US-0087759P.
PR	03-JUN-1998; 98US-0087827P.
PR	04-JUN-1998; 98US-0088021P.
PR	04-JUN-1998; 98US-0088025P.
PR	04-JUN-1998; 98US-0088028P.
PR	04-JUN-1998; 98US-0088029P.
PR	04-JUN-1998; 98US-0088030P.
PR	04-JUN-1998; 98US-0088033P.
PR	04-JUN-1998; 98US-0088326P.
PR	05-JUN-1998; 98US-0088167P.
PR	05-JUN-1998; 98US-0088202P.
PR	05-JUN-1998; 98US-0088212P.
PR	05-JUN-1998; 98US-0088217P.
PR	09-JUN-1998; 98US-0088655P.
PR	10-JUN-1998; 98US-0088722P.
PR	10-JUN-1998; 98US-0088730P.
PR	10-JUN-1998; 98US-0088734P.
PR	10-JUN-1998; 98US-0088738P.
PR	10-JUN-1998; 98US-0088740P.
PR	10-JUN-1998; 98US-0088741P.
PR	10-JUN-1998; 98US-0088742P.
PR	10-JUN-1998; 98US-0088810P.
PR	10-JUN-1998; 98US-0088811P.
PR	10-JUN-1998; 98US-0088824P.
PR	10-JUN-1998; 98US-0088825P.

QY 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMFRGRTAVFADQVIVGNASLRLKNV 120
Db 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMFRGRTAVFADQVIVGNASLRLKNV 120
QY 121 QLTDAITYKCYIIITSKGKNANLEYKTGAFSMPEVNVVDYNASSETLRCCEAPRWPQPTVV 180
Db 121 QLTDAITYKCYIIITSKGKNANLEYKTGAFSMPEVNVVDYNASSETLRCCEAPRWPQPTVV 180
QY 181 WASQVDQGANFSEVSNSTSFELNSENVTMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
Db 181 WASQVDQGANFSEVSNSTSFELNSENVTMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
QY 241 TESIKRSHQLQLLNKASLCVSSFFFAISWALLPLSPYMLK 282
Db 241 TESIKRSHQLQLLNKASLCVSSFFFAISWALLPLSPYMLK 282

RESULT 2

AAB12557
ID AAB12557 standard; protein; 282 AA.
XX AC AAB12557;

DT 07-NOV-2000 (first entry)

XX Human ovarian carcinoma antigen O8E protein SEQ ID NO:393.

DE Human; ovarian carcinoma; ovarian cancer; therapy; diagnosis;
KW tumour antigen; identification; cytostatic; gene therapy; vaccine.

XX Homo sapiens.

XX WO200036107-A2.

XX 22-JUN-2000.

XX PF 17-DEC-1999; 99WO-US030270.

XX 17-DEC-1998; 98US-00215681.

PR 17-DEC-1998; 98US-00216003.

PR 23-JUN-1999; 99US-00338933.

PR 24-SEP-1999; 99US-00404879.

XX (CORI-) CORIXA CORP.

XX Mitcam JL, King GE, Algate PA, Frudakis TN;

XX WPI; 2000-431589/37.

XX Immunogenic portion of an ovarian carcinoma protein and the nucleic acid
encoding it, useful for the diagnosis, prevention and treatment of
PT cancer, preferably ovarian cancer.

XX Example 2; Page 207; 299pp; English.

XX The present invention describes an isolated polypeptide comprising an
immunogenic portion of an ovarian carcinoma protein (or its variants).
CC Ovarian carcinoma proteins, and polynucleotides encoding them, have
CC cytosolic activity and can be used in gene therapy and vaccines. Ovarian
CC carcinoma polypeptides, nucleic acids, antibodies and vaccines are useful
CC for the prevention, diagnosis and treatment of cancer, preferably ovarian
CC cancer. AAA69691 to AAA70077 and AAB12552 to AAB12557 represent human
CC ovarian carcinoma polynucleotides and proteins used in the
CC exemplification of the present invention

XX Sequence 282 AA;

Query Match 100.0%; Score 1431; DB 3; Length 282;

Best Local Similarity 100.0%; Pred. No. 5,6e-118;

Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MASIGQILFWSIIIIILAGAIALLIIFGFSGRHSITVTTVASAGNIGEDGILSCTPEP 60

Db 1 MASIGQILFWSIIIIILAGAIALLIIFGFSGRHSITVTTVASAGNIGEDGILSCTPEP 60
QY 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMFRGRTAVFADQVIVGNASLRLKNV 120
Db 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMFRGRTAVFADQVIVGNASLRLKNV 120
QY 121 QLTDAITYKCYIIITSKGKNANLEYKTGAFSMPEVNVVDYNASSETLRCCEAPRWPQPTVV 180
Db 121 QLTDAITYKCYIIITSKGKNANLEYKTGAFSMPEVNVVDYNASSETLRCCEAPRWPQPTVV 180
QY 181 WASQVDQGANFSEVSNSTSFELNSENVTMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
Db 181 WASQVDQGANFSEVSNSTSFELNSENVTMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
QY 241 TESIKRSHQLQLLNKASLCVSSFFFAISWALLPLSPYMLK 282
Db 241 TESIKRSHQLQLLNKASLCVSSFFFAISWALLPLSPYMLK 282

RESULT 3

AAU29132
ID AAU29132 standard; protein; 282 AA.
XX AC AAU29132;

XX 18-DEC-2001 (first entry)

XX Human PRO polypeptide sequence #109.

KW PRO polypeptide; mammal; tumour; cancer; human; cattle; horse; sheep;
KW dog; cat; pig; goat; rabbit; tumour necrosis factor alpha; TNF-alpha;
KW blood; chondrocyte cell; cell proliferation; cell differentiation; colon;
KW adrenal; lung; breast; prostate; rectum; cervix; liver; genetic disorder.

XX Homo sapiens.

XX WO200168848-A2.

XX 20-SEP-2001.

XX 28-FEB-2001; 2001WO-US006520.

PR 01-MAR-2000; 2000WO-US005601.

PR 02-MAR-2000; 2000WO-US005841.

PR 03-MAR-2000; 2000US-0187202P.

PR 06-MAR-2000; 2000US-0186968P.

PR 14-MAR-2000; 2000US-0189320P.

PR 15-MAR-2000; 2000WO-US006884.

PR 21-MAR-2000; 2000US-0190828P.

PR 21-MAR-2000; 2000US-0191007P.

PR 21-MAR-2000; 2000US-0191048P.

PR 28-MAR-2000; 2000US-0192655P.

PR 29-MAR-2000; 2000US-0193032P.

PR 30-MAR-2000; 2000WO-US008439.

PR 04-APR-2000; 2000US-0194449P.

PR 11-APR-2000; 2000US-0195975P.

PR 11-APR-2000; 2000US-0196000P.

PR 11-APR-2000; 2000US-0196187P.

PR 11-APR-2000; 2000US-0196690P.

PR 18-APR-2000; 2000US-0196820P.

PR 18-APR-2000; 2000US-0198121P.

PR 25-APR-2000; 2000US-0198585P.

PR 25-APR-2000; 2000US-0199397P.

PR 25-APR-2000; 2000US-0199550P.

PR 03-MAY-2000; 2000US-0201516P.

PR 17-MAY-2000; 2000WO-US013705.

PR 22-MAY-2000; 2000WO-US014042.

PR 30-MAY-2000; 2000WO-US014941.
 PR 02-JUN-2000; 2000WO-US015264.
 PR 05-JUN-2000; 2000US-0209832P.
 PR 28-JUL-2000; 2000WO-US020710.
 PR 22-AUG-2000; 2000US-00644848.
 PR 24-AUG-2000; 2000WO-US023328.
 PR 08-NOV-2000; 2000WO-US030952.
 PR 01-DEC-2000; 2000WO-US032678.
 PR 20-DEC-2000; 2000WO-US034956.
 XX (GETH) GENENTECH INC.
 PA Baker KP, Chen J, Desnoyers L, Goddard A, Godowski PJ, Gurney AL;
 XX Pan J, Smith V, Watanabe CK, Wood WI, Zhang Z;
 PI WPI; 2001-602746/68.
 DR N-PSDB; AAS46033.
 XX Novel nucleic acids encoding PRO polypeptides, used to diagnose the
 PT presence of tumors, such as prostate and breast tumors, in mammals and to
 PT screen for modulators of the compounds.
 XX Claim 11; Fig 218; 774pp; English.
 PS Sequences AAU29024-AAU29328 represent PRO polypeptides of the invention.
 XX The PRO polypeptides and their associated nucleic acids can be used to
 CC detect the presence of a tumor in a mammal by comparing the level of
 CC expression of a PRO polypeptide in a test sample of cells from the animal
 CC and a control sample of normal cells, whereby a higher level of
 CC expression in the test sample indicates the presence of a tumor in the
 CC mammal. Mammals include dogs, cats, cattle, horses, sheep, pigs, goats
 CC and rabbits but are preferably human. The polypeptides can be used to
 CC stimulate tumour necrosis factor (TNF) alpha release from human blood,
 CC when contacted with it. A specific polypeptide can be used to stimulate
 CC the proliferation or differentiation of chondrocyte cells. The PRO
 CC proteins can be used to determine the presence of tumours and also
 CC susceptibility to tumour development, particularly adrenal, lung, colon,
 CC breast, prostate, rectal, cervical, or liver tumours, in mammalian
 CC subjects. The oligonucleotide probes specific for the PRO nucleic acids
 CC can be used for genetic analysis of individuals with genetic disorders
 XX Sequence 282 AA;

Query Match 100.0%; Score 1431; DB 4; Length 282;
 Best Local Similarity 100.0%; Pred. No. 5.6e-118;
 Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 Qy 1 MASLGQILFWSIIISIIILAGAIALIIIGFISGRHSITVTVASAGNIGEDGILSCTFEP 60
 Db 1 MASLGQILFWSIIISIIILAGAIALIIIGFISGRHSITVTVASAGNIGEDGILSCTFEP 60
 Qy 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
 Db 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
 Qy 121 QLTDAGTYKCYIITSKGNANLEYKTGAFSPMEVNVNDYNASSETLRCEAPRFPQPTVV 180
 Db 121 QLTDAGTYKCYIITSKGNANLEYKTGAFSPMEVNVNDYNASSETLRCEAPRFPQPTVV 180
 Qy 181 WASQVDQGANFSEVNTSFLNSENVTMKVSVLYNVTINNTYSCMIENDIAKATGDIKV 240
 Db 181 WASQVDQGANFSEVNTSFLNSENVTMKVSVLYNVTINNTYSCMIENDIAKATGDIKV 240
 Qy 241 TESEIKRSHLQLLNKASLCVSSFFPAISWALLPLSPYLMK 282
 Db 241 TESEIKRSHLQLLNKASLCVSSFFPAISWALLPLSPYLMK 282

RESULT 4
 AAB87555
 ID AAB87555 standard; protein; 282 AA.
 XX
 AC AAB87555;

XX 15-MAY-2001 (first entry)
 DT Human PRO1291.
 DE Human; PRO protein; mapping.
 KW Homo sapiens.
 XX OS
 XX WO200116318-A2.
 PN 08-MAR-2001.
 PD 24-AUG-2000; 2000WO-US023328.
 PF 01-SEP-1999; 99WO-US020111.
 XX 15-SEP-1999; 99WO-US021090.
 PR 07-DEC-1999; 99US-0169495P.
 PR 09-DEC-1999; 99US-0170262P.
 PR 11-JAN-2000; 2000US-0175481P.
 PR 18-FEB-2000; 2000WO-US004341.
 PR 18-FEB-2000; 2000WO-US004342.
 PR 22-FEB-2000; 2000WO-US004414.
 PR 01-MAR-2000; 2000WO-US005601.
 PR 03-MAR-2000; 2000US-0187202P.
 PR 21-MAR-2000; 2000US-0191007P.
 PR 30-MAR-2000; 2000WO-US008439.
 PR 25-APR-2000; 2000US-0199397P.
 PR 22-MAY-2000; 2000WO-US014042.
 PR 05-JUN-2000; 2000US-0209832P.
 XX (GETH) GENENTECH INC.
 PA Baton DL, Filvaroff E, Gerritsen ME, Goddard A, Godowski PJ;
 XX Grimaldi CJ, Gurney AL, Watanabe CK, Wood WI;
 PI WPI; 2001-183260/18.
 XX N-PSDB; AAF92087.
 DR Eighty four nucleic acids encoding PRO polypeptides, useful in molecular
 PT biology, including use as hybridization probes, and in chromosome and
 PT gene mapping.
 PT Claim 12; Fig 60; 278pp; English.
 XX The present sequence is a human PRO polypeptide (secreted and
 CC transmembrane). The PRO protein, and PRO agonists, PRO antagonists or
 CC anti-PRO antibodies are useful for preparation of a medicament useful in
 CC the treatment of a condition which is responsive to the PRO protein,
 CC agonists, antagonists or anti-PRO antibodies. The PRO protein may also be
 CC employed as molecular weight markers for protein electrophoresis. The PRO
 CC coding sequence has applications in molecular biology, including use as
 CC hybridisation probes, and in chromosome and gene mapping
 XX Sequence 282 AA;
 Query Match 100.0%; Score 1431; DB 4; Length 282;
 Best Local Similarity 100.0%; Pred. No. 5.6e-118;
 Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 Qy 1 MASLGQILFWSIIISIIILAGAIALIIIGFISGRHSITVTVASAGNIGEDGILSCTFEP 60
 Db 1 MASLGQILFWSIIISIIILAGAIALIIIGFISGRHSITVTVASAGNIGEDGILSCTFEP 60
 Qy 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
 Db 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
 Qy 121 QLTDAGTYKCYIITSKGNANLEYKTGAFSPMEVNVNDYNASSETLRCEAPRFPQPTVV 180
 Db 121 QLTDAGTYKCYIITSKGNANLEYKTGAFSPMEVNVNDYNASSETLRCEAPRFPQPTVV 180
 Qy 181 WASQVDQGANFSEVNTSFLNSENVTMKVSVLYNVTINNTYSCMIENDIAKATGDIKV 240

```
Db 181 WASQVDOGANFSEVSNSTFELNSENVTKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
Qy 241 TESIERSHQLNLNSKASLCVSSFFAISWALLPLSPYLMLK 282
Db 241 TESIERSHQLNLNSKASLCVSSFFAISWALLPLSPYLMLK 282

RESULT 5
AAB99204
ID AAB99204 standard; protein; 282 AA.
XX
AC AAB99204;
XX
DT 04-SEP-2001 (first entry)
XX
DE Human ovarian tumour-derived antigen O8E #1.
XX
KW Cytostatic; human; breast tumour protein; breast cancer; ovarian tumour;
KW antigen; O8E.
XX
OS Homo sapiens.
XX
PN WO200140269-A2.
XX
PD 07-JUN-2001.
XX
PF 29-NOV-2000; 2000WO-US032520.
XX
PR 30-NOV-1999; 99US-00451651.
PR 22-FEB-2000; 2000US-00510662.
PR 10-MAR-2000; 2000US-00523586.
PR 07-APR-2000; 2000US-00545068.
PR 15-MAY-2000; 2000US-00571025.
XX
PA (CORI-) CORIXA CORP.
XX
PI Dillon DC, Day CH, Jiang Y, Houghton RL, Mitcham JL, Wang A;
XX
DR WPI; 2001-356154/37.
XX
DR N-PSDB; AAH55681.
XX
PT Breast tumor polypeptides and the nucleic acids that encode them, useful
PT for the prevention, diagnosis and treatment of breast cancer.
XX
PS Example 3; Page 190; 221pp; English.
XX
CC The present invention relates to human breast tumour protein coding
CC sequences (see AAH55479-AAH55513, AAH55517-AAH55679 and AAH55682-
CC AAH55762). The breast tumour protein DNA sequences may be used in the
CC prevention, diagnosis and treatment of diseases associated with
CC inappropriate expression of the breast tumour protein e.g. breast cancer.
CC The present sequence is a human ovarian tumour-derived antigen, which was
CC used in an example from the present invention
XX
SQ Sequence 282 AA;

Query Match 100.0%; Score 1431; DB 4; Length 282;
Best Local Similarity 100.0%; Pred. No. 5.6e-118;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASLGQILFWSIIISIIILAGAIALLIIFGIGSRHSITVTTVASAGNIGEDGILSCTFEP 60
Db 1 MASLGQILFWSIIISIIILAGAIALLIIFGIGSRHSITVTTVASAGNIGEDGILSCTFEP 60

Qy 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMPRGRTAVFADQVIVGNASRLKNV 120
Db 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMPRGRTAVFADQVIVGNASRLKNV 120

Qy 121 QLTDAGTYKCYIITSKGNANLEYKTGAFSMPEVNVVDYNASSETLRCEAPRFPQPTVV 180
Db 121 QLTDAGTYKCYIITSKGNANLEYKTGAFSMPEVNVVDYNASSETLRCEAPRFPQPTVV 180
```

```
Qy 181 WASQVDOGANFSEVSNSTFELNSENVTKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
Db 181 WASQVDOGANFSEVSNSTFELNSENVTKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240

Qy 241 TESIERSHQLNLNSKASLCVSSFFAISWALLPLSPYLMLK 282
Db 241 TESIERSHQLNLNSKASLCVSSFFAISWALLPLSPYLMLK 282

RESULT 6
AAB65242
ID AAB65242 standard; protein; 282 AA.
XX
AC AAB65242;
XX
DT 02-APR-2001 (first entry)
XX
DE Human PRO1291 (UNQ659) protein sequence SEQ ID NO:291.
XX
KW Human; secreted and transmembrane protein; PRO; cytostatic; cell death;
KW cancer; chromosomal mapping; gene mapping; tissue typing;
XX
OS Homo sapiens.
XX
PN WO200073454-A1.
XX
PD 07-DEC-2000.
XX
PF 30-MAR-2000; 2000WO-US008439.
XX
PR 02-JUN-1999; 99WO-US012252.
PR 23-JUN-1999; 99US-0141037P.
PR 07-JUL-1999; 99US-0143048P.
PR 20-JUL-1999; 99US-0144758P.
PR 26-JUL-1999; 99US-0145698P.
PR 28-JUL-1999; 99US-0146222P.
PR 17-AUG-1999; 99US-0149396P.
PR 15-SEP-1999; 99WO-US021090.
PR 08-OCT-1999; 99WO-US021547.
PR 30-NOV-1999; 99US-0158663P.
PR 01-DEC-1999; 99WO-US028301.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 05-JAN-2000; 2000WO-US000219.
PR 06-JAN-2000; 2000WO-US000376.
PR 11-FEB-2000; 2000WO-US003565.
PR 18-FEB-2000; 2000WO-US004341.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US004914.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 15-MAR-2000; 2000WO-US006884.
PR 20-MAR-2000; 2000WO-US007377.
XX
PA (GETH ) GENENTECH INC.
XX
XX
PI Ashkenazi AJ, Baker KP, Borstein D, Desnoyers L, Eaton DL;
PI Ferrara N, Fong S, Gerber H, Gerritsen ME, Goddard A, Godowski PJ;
PI Grimaldi CJ, Gurney AL, Kijavini IJ, Napier MA, Pan J, Paoni NF;
PI Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM, Wood WI;
PI Zhang Z;
XX
XX
DR WPI; 2001-032160/04.
DR N-PSDB; AAF44205.
XX
XX
PT PRO polynucleotides used to produce polypeptides used to target bioactive
PT molecules such as toxins, radiolabels or antibodies, to specific cells,
PT to cause targeted cell death.
XX
XX
PS Claim 12; Fig 208; 935pp; English.
XX
```


CC The present invention describes human secreted and transmembrane PRO
 CC proteins. The PRO proteins have cytotstatic activity. The PRO proteins can
 CC be used for targeted delivery of bioactive molecules, such as toxins,
 CC radiolabels or antibodies, that cause cell death. PRO nucleotide
 CC sequences, and their fragments, can be used as hybridisation probes, in
 CC chromosomal and gene mapping, and in the generation of anti-sense RNA and
 CC DNA. They may also be used to produce transgenic animals which are used
 CC to develop and screen therapeutically useful reagents. The PRO nucleotide
 CC and protein sequence can be used for tissue typing and in treating
 CC cancer. Anti-PRO antibodies can be used in diagnostic assays. AAF44270 to
 CC AAF44470 represent PCR primers and hybridisation probes used in the
 CC isolation of human PRO sequences. AAF44087 to AAF44269 and AAB65154 to
 CC AAB65300 represent human PRO polynucleotide and protein sequences given
 CC in the exemplification of the present invention
 CC
 XX

SQ Sequence 282 AA;

Query Match 100.0%; Score 1431; DB 4; Length 282;
 Best Local Similarity 100.0%; Pred. No. 5.6e-118;
 Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASLGQLFWSIIISIIILAGATIALIIGFGISGRHSITVTVASAGNIGDGLSCTFEP 60
 Db |||||
 Qy 1 MASLGQLFWSIIISIIILAGATIALIIGFGISGRHSITVTVASAGNIGDGLSCTFEP 60
 Db |||||
 Qy 61 DIKLSDIVIOWLKEGVLGVHVEFKGKDELSEQDEMFRGTAVFADQVIVGNASRLKNV 120
 Db |||||
 Qy 61 DIKLSDIVIOWLKEGVLGVHVEFKGKDELSEQDEMFRGTAVFADQVIVGNASRLKNV 120
 Db |||||
 Qy 121 QLTDAGTYKCYIITSKGNANLEYKTFGAFSMPEVNVYDYNASSFTLRCEAPRFPQPTVV 180
 Db |||||
 Qy 121 QLTDAGTYKCYIITSKGNANLEYKTFGAFSMPEVNVYDYNASSFTLRCEAPRFPQPTVV 180
 Db |||||
 Qy 181 WASQVDOGANFSEVSNVTSFELNSENVMTKVVSVLYNVTINNTYSCMIENDIAKATGDIKV 240
 Db |||||
 Qy 181 WASQVDOGANFSEVSNVTSFELNSENVMTKVVSVLYNVTINNTYSCMIENDIAKATGDIKV 240
 Db |||||
 Qy 241 TESEIKRRSHLQLNLSKASLCVSSFFAISWALLPLSPYLMLK 282
 Db |||||
 Qy 241 TESEIKRRSHLQLNLSKASLCVSSFFAISWALLPLSPYLMLK 282
 Db |||||

RESULT 7

AAE20311

ID AAE20311 standard; protein; 282 AA.

AC AAE20311;

XX 18-JUN-2002 (first entry)

XX Human B7-H8 protein #1.

XX Human; B7-like protein; inflammation; tissue damage; immune disorder;
 KW Addison's disease; autoimmune haemolytic anaemia; autoimmune thyroiditis;
 KW diabetes mellitus; Crohn's disease; multiple sclerosis; allergy; cancer;
 KW rheumatoid arthritis; cardiovascular disorder; nervous system disorder;
 KW myocardial ischaemia; ulcerative colitis; reproductive system disorder;
 KW Alzheimer's disease; Parkinson's disease; endocrine disorder; hepatitis;
 KW diabetes mellitus; Grave's disease; Paget's disease; liver disorder;
 KW gastrointestinal disorder; irritable bowel syndrome; cerebral anoxia;
 KW dysphagia; hepatomegaly; neurological disease; infectious disease;
 KW epilepsy; gene therapy; B7-H8 protein; chromosome 1.

XX Homo sapiens.

XX Key Location/Qualifiers
 FH Peptide 1..24
 FT /label= Signal_peptide
 FT Protein 25..282
 FT /note= "Mature B7-H8 protein"

XX WO200202587-A1.

XX

PD 10-JAN-2002.

XX 29-JUN-2001; 2001WO-US020917.

XX 30-JUN-2000; 2000US-0215135P.

PR 14-AUG-2000; 2000US-0225266P.

XX (HUMA-) HUMAN GENOME SCI INC.

XX Fiscella M, Ni J, Ruben SM;

XX WPI; 2002-257198/30.

DR N-PSDB; AAD32519.

XX Isolated nucleic acids encoding human B7-like polypeptides, useful for

PT diagnosis and treatment of e.g. inflammation, cancer, immune disorders

PT such as Addison's disease, and cardiovascular disorders such as

PT myocardial ischemias.

XX Example 1; Fig 1; 493pp; English.

XX The present invention relates to novel human B7-like polypeptides and

CC polynucleotides encoding such proteins. Sequences of the invention are

CC used for preventing, treating or ameliorating a medical condition in a

CC mammalian subject. The polynucleotides and polypeptides are administered

CC to subjects having a disorder related to B-7 Like polypeptides, such as

CC inappropriate or excessive inflammation which can lead to tissue damage

CC or even death, where the inflammation is brought about by the activation

CC of certain cells in the body e.g. T cells and may involve disorders

CC related to immune system. The nucleic acids, proteins, antibodies,

CC agonists and antagonists of the invention are useful in the diagnosis,

CC treatment and prevention of cancer (e.g. cancers of the adrenal gland,

CC bone, bone marrow, breast, gastrointestinal tract, liver, urogenital or

CC lung), immune disorders (e.g., Addison's disease, allergies, autoimmune

CC haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's

CC disease, multiple sclerosis, rheumatoid arthritis, ulcerative colitis),

CC cardiovascular disorders (e.g., myocardial ischaemias), nervous system

CC disorders (Alzheimer's disease, Parkinson's disease), endocrine disorders

CC (e.g., diabetes mellitus, Grave's disease), reproductive system disorders

CC (e.g., cryptorchism, Paget's disease), gastrointestinal disorders (e.g.,

CC dysphagia, irritable bowel syndrome), liver disorders (e.g., hepatitis,

CC hepatomegaly), neurological diseases (e.g., cerebral anoxia and epilepsy)

CC and infectious diseases such as viral, bacterial, fungal and parasitic

CC infections. Sequences of the invention are also used in gene therapy. The

CC present sequence is human B7-H8 protein. B7-H8 gene is located on

CC chromosome 1

XX

SQ Sequence 282 AA;

Query Match 100.0%; Score 1431; DB 5; Length 282;

Best Local Similarity 100.0%; Pred. No. 5.6e-118;

Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASLGQLFWSIIISIIILAGATIALIIGFGISGRHSITVTVASAGNIGDGLSCTFEP 60

Db |||||

1 MASLGQLFWSIIISIIILAGATIALIIGFGISGRHSITVTVASAGNIGDGLSCTFEP 60

Qy 61 DIKLSDIVIOWLKEGVLGVHVEFKGKDELSEQDEMFRGTAVFADQVIVGNASRLKNV 120

Db |||||

61 DIKLSDIVIOWLKEGVLGVHVEFKGKDELSEQDEMFRGTAVFADQVIVGNASRLKNV 120

Qy 121 QLTDAGTYKCYIITSKGNANLEYKTFGAFSMPEVNVYDYNASSFTLRCEAPRFPQPTVV 180

Db |||||

121 QLTDAGTYKCYIITSKGNANLEYKTFGAFSMPEVNVYDYNASSFTLRCEAPRFPQPTVV 180

Qy 181 WASQVDOGANFSEVSNVTSFELNSENVMTKVVSVLYNVTINNTYSCMIENDIAKATGDIKV 240

Db |||||

181 WASQVDOGANFSEVSNVTSFELNSENVMTKVVSVLYNVTINNTYSCMIENDIAKATGDIKV 240

Qy 241 TESEIKRRSHLQLNLSKASLCVSSFFAISWALLPLSPYLMLK 282

Db |||||

241 TESEIKRRSHLQLNLSKASLCVSSFFAISWALLPLSPYLMLK 282

Qy

Db

RESULT 8
 ABG96445
 ID ABG96445 standard; protein; 282 AA.
 XX
 AC ABG96445;
 XX
 DT 11-DEC-2002 (first entry)
 XX
 DE Human ovarian cancer marker OV88.
 XX
 KW Human; ovarian cancer; marker; cancer; familial history; brain disorder;
 KW central nervous system disorder; bacterial meningitis; viral meningitis;
 KW Alzheimer's disease; Parkinson's disease; cerebral oedema; hydrocephalus;
 KW brain herniation; inflammation; encephalitis; testicular disorder;
 KW nontuberculous granulomatous orchitis; connective tissue disorder;
 KW heart disorder; ischaemic heart disease; atherosclerosis; neoplasm;
 KW histological type; carcinogenic; ovarian cancer marker.
 XX
 OS Homo sapiens.
 XX
 PN WO200271928-A2.
 XX
 PD 19-SEP-2002.
 XX
 PF 14-MAR-2002; 2002WO-US007826.
 XX
 PR 14-MAR-2001; 2001US-0276025P.
 PR 14-MAR-2001; 2001US-0276026P.
 PR 10-AUG-2001; 2001US-0311732P.
 PR 19-SEP-2001; 2001US-0323580P.
 PR 26-SEP-2001; 2001US-0324967P.
 PR 26-SEP-2001; 2001US-0325102P.
 PR 26-SEP-2001; 2001US-0325149P.
 XX
 PA (MILL-) MILLENNIUM PHARM INC.
 XX
 PI Monahan JB, Gannavarapu M, Hoersch S, Kamatkar S, Kovatis SG;
 PI Meyers RE, Morrissey MF, Olandt PJ, Sen A, Vieby PO, Mills GB;
 PI Baat RC, Lu K, Schmandt RE, Zhao X, Glatt K;
 XX
 DR WPI; 2002-723277/78.
 DR N-PSDB; ABS76544.
 XX
 PT Assessing whether a patient is afflicted with ovarian cancer, useful in
 PT assessing the stage or progression of the disease, comprises comparing
 PT the expression level of a cancer marker in a sample from a patient and
 PT from a non cancer patient.
 XX
 PS Disclosure; Page 468-469; 481pp; English.
 XX
 CC The present invention relates to a new method for assessing whether a
 CC patient is afflicted with ovarian cancer. The method involves comparing
 CC the expression level of a marker in a patient sample and the normal level
 CC of expression of the marker in a control non-ovarian cancer sample, where
 CC the marker is selected from 363 cancer markers described in the
 CC specification. The method of the invention is useful in diagnosing or
 CC characterising cancer, in detecting the presence of cancer as early as
 CC possible, and the recurrence of ovarian cancer. The method may also be of
 CC particular use with patients having an enhanced risk of developing
 CC ovarian cancer (e.g. patients having a familial history of ovarian
 CC cancer). The cancer markers may be used in the management and treatment
 CC of e.g. brain and central nervous system disorders (e.g. bacterial and
 CC viral meningitis, Alzheimer's disease or Parkinson's disease), brain
 CC disorders (e.g. cerebral oedema, hydrocephalus or brain herniations),
 CC inflammations (e.g. bacterial or viral meningitis or encephalitis),
 CC testicular disorders (e.g. nontuberculous granulomatous orchitis),
 CC connective tissue disorders, or heart disorders (e.g. ischaemic heart
 CC disease or atherosclerosis). The compositions and methods may also be
 CC used in assessing the histological type of neoplasm associated with
 CC ovarian cancer, monitoring the progression of ovarian cancer, determining
 CC whether ovarian cancer has metastasized or is likely to metastasize,
 CC selecting a composition for inhibiting ovarian cancer, assessing the

CC ovarian carcinogenic potential of a compound, or inhibiting ovarian
 CC cancer or at risk of developing ovarian cancer. The present amino acid
 CC sequence represents one of the ovarian cancer markers described in the
 CC invention
 XX
 SQ Sequence 282 AA;
 XX
 Query Match 100.0%; Score 1431; DB 5; Length 282;
 Best Local Similarity 100.0%; Pred. No. 5.6e-118;
 Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 Qy 1 MASIGQILFWISIIIIILAGAIALLIIFGIGSRHSITVTTVASAGNIGEDGILSCTPEP 60
 Db 1 MASIGQILFWISIIIIILAGAIALLIIFGIGSRHSITVTTVASAGNIGEDGILSCTPEP 60
 Qy 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMFERGRTAVFADQVIVGNASLRLKNV 120
 Db 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMFERGRTAVFADQVIVGNASLRLKNV 120
 Qy 121 QLTDAGTYKCYIITSKGGNANLEYKTGAFSMPENVVDYNASSETLRCEAPRFPQPTVW 180
 Db 121 QLTDAGTYKCYIITSKGGNANLEYKTGAFSMPENVVDYNASSETLRCEAPRFPQPTVW 180
 Qy 181 WASQVDOGANFSEVSNTPFELNSENVTMKVSVLYNVTINNTYSCMIENDIAKATGDIKV 240
 Db 181 WASQVDOGANFSEVSNTPFELNSENVTMKVSVLYNVTINNTYSCMIENDIAKATGDIKV 240
 Qy 241 TESEIKRSHLQLLNKASLCVSSFFFAISWALLPLSPYLMLK 282
 Db 241 TESEIKRSHLQLLNKASLCVSSFFFAISWALLPLSPYLMLK 282
 RESULT 9
 AAU77766
 ID AAU77766 standard; protein; 282 AA.
 XX
 AC AAU77766;
 XX
 DT 05-JUN-2002 (first entry)
 XX
 DE Tumour associated antigenic target polypeptide (TAT) 136.
 XX
 KW Tumour associated antigenic target polypeptide; TAT; cancer;
 KW breast cancer; colorectal cancer; lung cancer; ovarian cancer;
 KW central nervous system cancer; liver cancer; bladder cancer;
 KW pancreatic cancer; cervical cancer; melanoma; leukaemia; TAT136.
 XX
 OS Homo sapiens.
 XX
 FH Key Location/Qualifiers
 FT Peptide 1..28
 FT Protein /label= Signal_peptide
 FT /label= Mature_TAT136
 FT /note= "Tumour associated antigenic target polypeptide"
 FT Region 52..58
 FT /label= N-myristoylation_site
 FT Region 112..116
 FT /label= N-glycosylation_site
 FT Domain 119..123
 FT /label= Immunoglobulin_domain
 FT Region 126..132
 FT /label= N-myristoylation_site
 FT Region 160..164
 FT /label= N-glycosylation_site
 FT Region 188..194
 FT /label= N-myristoylation_site
 FT Region 190..194
 FT /label= N-glycosylation_site
 FT Region 196..200
 FT /label= N-glycosylation_site
 FT Region 205..209
 FT /label= N-glycosylation_site

(ANGE-) AMGEN INC.
 Fox M, Sullivan JK, Fang M;
 WPI; 2002-171639/22.
 N-PSDB; AAD29253.
 Novel B7-like polypeptides, polynucleotides and their modulators useful
 for prevention and treatment of reproductive, immune and proliferative
 disorders, e.g. cancer, arteriosclerosis.
 Claim 13; Fig 1A-1B; 133pp; English.
 The present invention relates to an isolated B7-like (B7-L) polypeptide
 and its polynucleotide. B7-1 and its modulators are useful for treating
 reproductive disorders (e.g. infertility, miscarriage, preterm labour and
 delivery and endometriosis) and proliferative disorders. Antibodies,
 soluble proteins comprising extracellular domains and other regulators of
 B7-L are useful for enhancing the immune response to tumours. B7-1 plays
 a role in growth and maintenance of cancer cells based on the observation
 of seminal vesicle hyperplasia in transgenic mice overexpressing B7-1.
 Modulators of B7-1 are useful for the treatment of cancer e.g. seminal
 vesicle, lung, brain, breast, ovarian, testicular cancer and cancers of
 haematopoietic system. B7-1 and their modulators are useful to treat
 autoimmune diseases such as systemic lupus erythematosus, rheumatoid
 arthritis, immune thrombocytopenic purpura and psoriasis, chronic
 inflammatory disease such as inflammatory bowel disease (Crohn's disease
 and ulcerative colitis), Grave's disease, Hashimoto's thyroiditis and
 diabetes mellitus. They are also useful as immunosuppressive agents for
 bone marrow and organ transplantation or to prolong graft survival.
 Modulators of B7-L are also useful for diagnosis and treatment of
 diseases involving abnormal cell proliferation, arteriosclerosis and
 vascular restenosis. Soluble B7-L serves as vaccine adjuvants.
 Antagonists of B7-L are useful for alleviation of toxic shock syndrome or
 allosterisation due to blood transfusions, and for treatment of
 multiple sclerosis, allergy, asthma and hypersensitivity reactions,
 nephropathies (e.g. glomerulonephritis), skin disorders (pemphigus,
 pemphigoid), endocrinopathies, various pneumopathies, vasculopathies,
 coeliac disease, anaemia, thrombocytopenia, Guillain-Barre syndrome and
 myasthenia gravis, and lymphoproliferative disorders such as multiple
 myeloma. B7-L gene is useful in gene therapy and to map the locations of
 B7-L gene and related genes on chromosomes, as hybridisation probes in
 diagnostic assays, for isolating corresponding chromosomal B7-L genes,
 and to identify heritable tissue-degenerating diseases. The present
 sequence is human B7-L protein

Query Match 100.0%; Score 1431; DB 5; Length 282;
 Best Local Similarity 100.0%; Pred. No. 5.6e-118;
 Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASLGQLFWSIIISIIILAGAILIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60
 Db 1 MASLGQLFWSIIISIIILAGAILIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60

Qy 61 DIKLSDIVIQWLKEGVGLVHEFKEGKDELSEQDEMFRGTAVFADQVIVGNASLRKNV 120
 Db 61 DIKLSDIVIQWLKEGVGLVHEFKEGKDELSEQDEMFRGTAVFADQVIVGNASLRKNV 120

Qy 121 QLTDAGTYKCYIITSKGNANLEYKKGAFSMPBVVDYNASSFTLCEAPRFPQPTVV 180
 Db 121 QLTDAGTYKCYIITSKGNANLEYKKGAFSMPBVVDYNASSFTLCEAPRFPQPTVV 180

Qy 181 WASQVDCGANFSEVNSVTSFELNSENVTKVSVLYNVTINNTYSCMIENDIAKATGDIKV 240
 Db 181 WASQVDCGANFSEVNSVTSFELNSENVTKVSVLYNVTINNTYSCMIENDIAKATGDIKV 240

Qy 241 TESIERSHLOLNSKASLCVSSFFAISWALLPLSPYMLK 282
 Db 241 TESIERSHLOLNSKASLCVSSFFAISWALLPLSPYMLK 282

RESULT 15

ABB09879
 ID ABB09879 standard; protein; 282 AA.
 XX
 AC ABB09879;
 XX
 DT 30-JUL-2002 (first entry)
 XX
 DE Amino acid sequence of the OREO gene (gene B).
 XX
 KW Human; gene A; ovarian tumour; gene B; OREO; ovarian cancer.
 XX
 OS Homo sapiens.
 XX
 FH Key Location/Qualifiers
 XX Domain 12..31
 FT /note= "predicted transmembrane domain"
 FT Domain 46..145
 FT /note= "predicted Ig domain"
 FT Modified-site 112
 FT /note= "N-glycosylation site"
 FT Modified-site 160
 FT /note= "N-glycosylation site"
 FT Modified-site 190
 FT /note= "N-glycosylation site"
 FT Modified-site 196
 FT /note= "N-glycosylation site"
 FT Modified-site 205
 FT /note= "N-glycosylation site"
 FT Modified-site 216
 FT /note= "N-glycosylation site"
 FT Modified-site 220
 FT /note= "N-glycosylation site"
 XX
 W0200194641-A2.
 PN
 PD 13-DEC-2001.
 XX
 PD 11-JUN-2001; 2001WO-US018700.
 XX
 PR 09-JUN-2000; 2000US-0210451P.
 XX
 PA (IDEC-) IDEC PHARM CORP.
 XX
 PI Ople B, McLachlan K, Heard C;
 XX
 DR WPI; 2002-404365/43.
 DR N-PSDB; ABL56582.
 XX
 PT New polynucleotide and corresponding antigens from human ovarian cancer
 FT cells, useful for treatment and diagnosis of ovarian cancer.
 XX
 PS Claim 12; Fig 7b; 71pp; English.
 XX
 CC The present sequence represents a protein designated OREO. The OREO (Ople
 CC RBA of Epithelial Tissue vs. Ovary tumour) gene is a novel gene, also
 CC designated gene B. This gene was identified by representational
 CC difference analysis (RDA) screening, and is selectively expressed by
 CC certain human ovarian tumours. The specification also describes gene A,
 CC identified by the same method. Gene A and B polynucleotides are useful
 CC for detecting ovarian cancer. Their polypeptides are useful for treating
 CC ovarian cancer
 XX
 SQ Sequence 282 AA;
 Query Match 100.0%; Score 1431; DB 5; Length 282;
 Best Local Similarity 100.0%; Pred. No. 5.6e-118;
 Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASLGQLFWSIIISIIILAGAILIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60
 Db 1 MASLGQLFWSIIISIIILAGAILIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60

Qy	61	DIKLSDIVIQWLKEGVLGLVHEPKEGKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV	120
Db	61		
Qy	121	DIKLSDIVIQWLKEGVLGLVHEPKEGKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV	120
Db	121		
Qy	181	QLTDAGTYKCYIITSKKGKGNANLEYKTGAFSMPENVVDYNASSETLRCEAPRWFPPQPTVV	180
Db	181		
Qy	241	QLTDAGTYKCYIITSKKGKGNANLEYKTGAFSMPENVVDYNASSETLRCEAPRWFPPQPTVV	180
Db	241		
Qy	282	WASQVDCGANFSEVSNTSFELNSENVTMKVSVLYNVVTINNTYSCMIENDIAKATGDIKV	240
Db	282		
Qy	282	WASQVDCGANFSEVSNTSFELNSENVTMKVSVLYNVVTINNTYSCMIENDIAKATGDIKV	240
Db	282		
Qy	282	TESEIKRSHLOLLNSKASLCVSSFFAISWALLPLSPYLMK	282
Db	282		
Qy	282	TESEIKRSHLOLLNSKASLCVSSFFAISWALLPLSPYLMK	282
Db	282		

Search completed: April 19, 2005, 07:13:28
Job time : 98 secs

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	Query Match	100.0%;	Score 282;	DB 9;	Length 282;
	Best Local Similarity	100.0%;	Pred. No. 2e-263;		
	Matches 282;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0;
QY	1	MASLGQILFWISIISIIIIAGATALLIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP	60		
Db	1	MASLGQILFWISIISIIIIAGATALLIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP	60		
QY	61	DIKLSDIVTQWLKEGVGLGVHPFKGKDELSEQDEMPFRGRTAVPADQVIVGNASRLKNV	120		
Db	61	DIKLSDIVTQWLKEGVGLGVHPFKGKDELSEQDEMPFRGRTAVPADQVIVGNASRLKNV	120		

Qy 121 QLTDAAGTYKCYIITSKGNANLEYKTFGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180
Db 121 QLTDAAGTYKCYIITSKGNANLEYKTFGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180
Qy 181 WASQVDGAFSEVSNSTSFELNSENVMTKVVSVLYNVTINNTYSCMIENDIAKATGDIKV 240
Db 181 WASQVDGAFSEVSNSTSFELNSENVMTKVVSVLYNVTINNTYSCMIENDIAKATGDIKV 240
Qy 241 TESEIKRRSHLQLNSKASLCVSSFFAISWALLPLSPYMLK 282
Db 241 TESEIKRRSHLQLNSKASLCVSSFFAISWALLPLSPYMLK 282

RESULT 2

US-09-850-178-33
; Sequence 33, Application US/09850178
; Patent No. US20020034749A1
; GENERAL INFORMATION:
; APPLICANT: Abbott Laboratories
; APPLICANT: Billing-Medel, Patricia A.
; APPLICANT: Cohen, Maurice
; APPLICANT: Colipitts, Tracey L.
; APPLICANT: Friedman, Paula N.
; APPLICANT: Russell, John C.
; APPLICANT: Granados, Edward N.
; APPLICANT: Hodges, Steven C.
; APPLICANT: Klass, Michael R.
; APPLICANT: Kratochvil, Jon D.
; APPLICANT: Roberts-Rapp, Lisa
; APPLICANT: Stroupe, Stephen D.
; APPLICANT: Gordon, Julian
; TITLE OF INVENTION: REAGENTS AND METHODS USEFUL FOR
; FILE OF INVENTION: DETECTING DISEASES OF THE BREAST
; FILE REFERENCE: 6251.US.P1
; CURRENT APPLICATION NUMBER: US/09/850.178
; CURRENT FILING DATE: 2001-05-07
; PRIOR APPLICATION NUMBER: US 08/972,376
; PRIOR FILING DATE: 1997-11-18
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: Fast-Seq for Windows Version 4.0
; SEQ ID NO 33
; LENGTH: 282
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-850-178-33

Query Match 100.0%; Score 282; DB 9; Length 282;
Best Local Similarity 100.0%; Pred. No. 2e-263;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 MASLGQILFWSIISIIILAGAILIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60
Db 1 MASLGQILFWSIISIIILAGAILIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60
Qy 61 DIKLSDIVIOWLKEGVLGVHFEKGEKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
Db 61 DIKLSDIVIOWLKEGVLGVHFEKGEKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
Qy 121 QLTDAAGTYKCYIITSKGNANLEYKTFGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180
Db 121 QLTDAAGTYKCYIITSKGNANLEYKTFGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180
Qy 181 WASQVDGAFSEVSNSTSFELNSENVMTKVVSVLYNVTINNTYSCMIENDIAKATGDIKV 240
Db 181 WASQVDGAFSEVSNSTSFELNSENVMTKVVSVLYNVTINNTYSCMIENDIAKATGDIKV 240
Qy 241 TESEIKRRSHLQLNSKASLCVSSFFAISWALLPLSPYMLK 282
Db 241 TESEIKRRSHLQLNSKASLCVSSFFAISWALLPLSPYMLK 282

RESULT 3

US-09-877-065-8
; Sequence 8, Application US/09877065
; Patent No. US20020051990A1
; GENERAL INFORMATION:
; APPLICANT: OPLE, ERIC
; APPLICANT: McLachlan, Karen
; APPLICANT: Heard, Cheryl J.
; TITLE OF INVENTION: NOVEL GENE TARGETS AND LIGANDS THAT BIND THERETO FOR
; TITLE OF INVENTION: TREATMENT AND DIAGNOSIS OF OVARIAN CARCINOMAS
; FILE REFERENCE: 037003-0280631
; CURRENT APPLICATION NUMBER: US/09/877,065
; CURRENT FILING DATE: 2001-06-11
; PRIOR APPLICATION NUMBER: 60/210,451
; PRIOR FILING DATE: 2000-06-09
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 8
; LENGTH: 282
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-877-065-8

Query Match 100.0%; Score 282; DB 9; Length 282;
Best Local Similarity 100.0%; Pred. No. 2e-263;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 MASLGQILFWSIISIIILAGAILIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60
Db 1 MASLGQILFWSIISIIILAGAILIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60
Qy 61 DIKLSDIVIOWLKEGVLGVHFEKGEKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
Db 61 DIKLSDIVIOWLKEGVLGVHFEKGEKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
Qy 121 QLTDAAGTYKCYIITSKGNANLEYKTFGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180
Db 121 QLTDAAGTYKCYIITSKGNANLEYKTFGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180
Qy 181 WASQVDGAFSEVSNSTSFELNSENVMTKVVSVLYNVTINNTYSCMIENDIAKATGDIKV 240
Db 181 WASQVDGAFSEVSNSTSFELNSENVMTKVVSVLYNVTINNTYSCMIENDIAKATGDIKV 240
Qy 241 TESEIKRRSHLQLNSKASLCVSSFFAISWALLPLSPYMLK 282
Db 241 TESEIKRRSHLQLNSKASLCVSSFFAISWALLPLSPYMLK 282

RESULT 4

US-09-989-722-291
; Sequence 291, Application US/09989722
; Patent No. US20020072067A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Baton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Napier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.

APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William I.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C63
CURRENT APPLICATION NUMBER: US/09/989,722
CURRENT FILING DATE: 2001-11-19
PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/065186
PRIOR FILING DATE: 1997-11-12
PRIOR APPLICATION NUMBER: 60/065311
PRIOR FILING DATE: 1997-11-13
PRIOR APPLICATION NUMBER: 60/066770
PRIOR FILING DATE: 1997-11-24
PRIOR APPLICATION NUMBER: 60/075945
PRIOR FILING DATE: 1998-02-25
PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/083322
PRIOR FILING DATE: 1998-04-28
PRIOR APPLICATION NUMBER: 60/084600
PRIOR FILING DATE: 1998-05-07
PRIOR APPLICATION NUMBER: 60/087106
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; PRIOR APPLICATION NUMBER: 60/091544
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; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091633
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091978
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 282; DB 9; Length 282;

Best Local Similarity 100.0%; Pred. No. 2e-263; Mismatches 0; Indels 0; Gaps 0;
Matches 282; Conservative 0;

Qy	1	MASIGQLFWSIISIIILAGALAIIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP	60
Db	1	MASIGQLFWSIISIIILAGALAIIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP	60
Qy	61	DIKLSDIVIOWLKEGVGLVHFEKGEKDELSEQDEMFRGTAVPADQVIVGNASLRKNV	120
Db	61	DIKLSDIVIOWLKEGVGLVHFEKGEKDELSEQDEMFRGTAVPADQVIVGNASLRKNV	120
Qy	121	QLTDAGTYKCYIITSKGNANLEYKTFGAFSMEPVNVVDYNASSETLRCEAPRFPQPTVV	180
Db	121	QLTDAGTYKCYIITSKGNANLEYKTFGAFSMEPVNVVDYNASSETLRCEAPRFPQPTVV	180
Qy	181	WASQVDCGANFSEVSNSTFELNSENVMTKVVSVLYNVTINNTYSCTMIENDIAKATGDIKV	240
Db	181	WASQVDCGANFSEVSNSTFELNSENVMTKVVSVLYNVTINNTYSCTMIENDIAKATGDIKV	240
Qy	241	TESEIKRSHQLLNKSKASLCVSSFFAISWALLPLSPYMLK	282
Db	241	TESEIKRSHQLLNKSKASLCVSSFFAISWALLPLSPYMLK	282

RESULT 5

US-09-989-723-291
; Sequence 291, Application US/09989723
; Patent No. US20020072092A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher

; APPLICANT: Gurney, Austin L.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Nepier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: P2730P162
; CURRENT APPLICATION NUMBER: US/09/989, 723
; CURRENT FILING DATE: 2001-11-19
; PRIOR APPLICATION NUMBER: 60/049787
; PRIOR FILING DATE: 1997-06-16
; PRIOR APPLICATION NUMBER: 60/062250
; PRIOR FILING DATE: 1997-10-17
; PRIOR APPLICATION NUMBER: 60/065186
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; PRIOR FILING DATE: 1998-02-25
; PRIOR APPLICATION NUMBER: 60/078910
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; PRIOR FILING DATE: 1998-06-24
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; PRIOR FILING DATE: 1998-06-24
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; PRIOR FILING DATE: 1998-06-24
; PRIOR APPLICATION NUMBER: 60/090557
; PRIOR FILING DATE: 1998-06-24
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; PRIOR APPLICATION NUMBER: 60/091360
; PRIOR FILING DATE: 1998-07-01
; PRIOR APPLICATION NUMBER: 60/091478
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091544
; PRIOR FILING DATE: 1998-07-01
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; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091626
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091633
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091978
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 282; DB 9; Length:282;
Best Local Similarity 100.0%; Pred. No. 28-263; Indels 0; Gaps 0;
Matches 282; Conservative 0; Mismatches 0;

Qy 1 MASLQILFWSIISIIILAGAILIIGFISGRHSITVTTVASAGNIGDGLSCTFEP 60
Db 1 MASLQILFWSIISIIILAGAILIIGFISGRHSITVTTVASAGNIGDGLSCTFEP 60
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Db 61 DIKLSDIVIQLKEGVLGVHVEFKGKDELSQDENFRGRTAVFADQVIVGNASLRKNV 120
Qy 121 QLTDACTYKCIITTSKGNANLEYKTGAFSMEPVNDVYNASSETLRCEAPRFPQPTVV 180
Db 121 QLTDACTYKCIITTSKGNANLEYKTGAFSMEPVNDVYNASSETLRCEAPRFPQPTVV 180
Qy 181 WASQVDQGANFSEVSNSTSFELNSENVTMKVSVLYNVNTINNTYSCHMIENDIAKATGDIKV 240
Db 181 WASQVDQGANFSEVSNSTSFELNSENVTMKVSVLYNVNTINNTYSCHMIENDIAKATGDIKV 240
Qy 241 TESEIKRRSHLQLLNKSKSLCVSSFFAISWALLPLSPYMLK 282
Db 241 TESEIKRRSHLQLLNKSKSLCVSSFFAISWALLPLSPYMLK 282

RESULT 6
US-09-989-279-291
; Sequence 291, Application US/09989279
; Patent No. US20020072496A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David

APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Fong, Sherman
APPLICANT: Gerber, Hanspeter
APPLICANT: Geritsen, Mary E.
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Gurney, Austin L.
APPLICANT: Kijavini, Ivar J.
APPLICANT: Napier, Mary A.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Watanabe, Colin K.
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William I.
APPLICANT: Zhang, Zenin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE REFERENCE: P2730P1C56
CURRENT APPLICATION NUMBER: US/09/989,279
CURRENT FILING DATE: 2001-11-19
PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/062250
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;; PRIOR FILING DATE: 1998-07-07
;; PRIOR APPLICATION NUMBER: 60/092182
;; PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 282; DB 9; Length 282;
Best Local Similarity 100.0%; Pred. No. 2e-263;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db 1 MASLGQLFWFSIIIIILAGAIALLIGFGISGRHSITVTVVASAGNIGEDGILSCTFEP 60

Qy 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMFRGRTAVPADQVIVGNASRLKNV 120
Db 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMFRGRTAVPADQVIVGNASRLKNV 120

Qy 121 QLTDAGTYKCYIITSKGGNANLEYKTKGAFSMPEVNVVDYNASSETLACEAPRFPQPTVV 180
Db 121 QLTDAGTYKCYIITSKGGNANLEYKTKGAFSMPEVNVVDYNASSETLACEAPRFPQPTVV 180

Qy 181 WASQVDGAFSEVNSVTSFELNSENVTKVSVLVYNTVNTTYSMTENDIAKATGDIKV 240
Db 181 WASQVDGAFSEVNSVTSFELNSENVTKVSVLVYNTVNTTYSMTENDIAKATGDIKV 240

Qy 241 TESIKRSHLOLNSKASLCVSSFFAISWALLPLSPVLMK 282
Db 241 TESIKRSHLOLNSKASLCVSSFFAISWALLPLSPVLMK 282

RESULT 7

US-09-989-727-291
; Sequence 291, Application US/09989727
; Patent No. US20020072497A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
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; APPLICANT: Gerber, Hanspeter
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; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: P2730P1G65
; CURRENT APPLICATION NUMBER: US/09/989,727
; CURRENT FILING DATE: 2001-11-19
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; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

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Db      61 DIKLSDIVIQWLKEGVLGVHFEKCKDELSEODEMFRGRTAVFADQVIVGNASRLKNV 120
Oy      121 QLTDACTYKCYIITSKGNANLEVYKTGAFSPMPVNVNDYNASSETLRCEAPRFPPTVV 180
Db      121 QLTDACTYKCYIITSKGNANLEVYKTGAFSPMPVNVNDYNASSETLRCEAPRFPPTVV 180
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Qy 241 TSEIKRSHQLLNKSKASLCVSSFFAISWALLPLSPYMLK 282
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US-09-910-689-208
; Sequence 208, Application US/09910689
; Patent No. US20020081609A1
; GENERAL INFORMATION:
; APPLICANT: Dillon, Davin C.
; APPLICANT: Dav, Craig H.
; APPLICANT: Jiang, Yuciu
; APPLICANT: Houghton, Raymond L.
; APPLICANT: Mitcham, Jennifer
; APPLICANT: Wang, Tongtong
; APPLICANT: McNeill, Patricia D.
; APPLICANT: Harlocker, Susan L.
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
; TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
; FILE REFERENCE: 210121.491C6
; CURRENT APPLICATION NUMBER: US/09/910.689
; CURRENT FILING DATE: 2001-07-20
; NUMBER OF SEQ ID NOS: 307
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 208
; LENGTH: 282
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-910-689-208

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Qy 241 TSEIKRSHQLLNKSKASLCVSSFFAISWALLPLSPYMLK 282
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RESULT 9

US-09-989-731-291
; Sequence 291, Application US/09989731
; Patent No. US20020103125A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman

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; APPLICANT: Godowski, Paul J.
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; APPLICANT: Napier, Mary A.
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; APPLICANT: Paoni, Nicholas F.
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; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: P2730PIC70
; CURRENT APPLICATION NUMBER: US/09/989.731
; CURRENT FILING DATE: 2001-11-20
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; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 282; DB 9; Length 282;
Best Local Similarity 100.0%; Pred. No. 2e-263;

Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy	1	MASLGQLFWSIIISIIILAGAILIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP	60
Db	1	MASLGQLFWSIIISIIILAGAILIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP	60
Qy	61	DIKLSDIVIOWLKEGVLGVHFEKGEKDELSEODEMPRGRTAVFADQVIVGNASLRKNV	120
Db	61	DIKLSDIVIOWLKEGVLGVHFEKGEKDELSEODEMPRGRTAVFADQVIVGNASLRKNV	120
Qy	121	QLTDAGTYKCYIITSKGNANLEYKTGAFSMEPVNDVYNASSETLRCEAPRFPQPTVV	180
Db	121	QLTDAGTYKCYIITSKGNANLEYKTGAFSMEPVNDVYNASSETLRCEAPRFPQPTVV	180
Qy	181	WASQVDOGANFSEVSNTSFELNSENVTMKVSVLYNVNTINNTYSCEMIENDIAKATGDIKV	240
Db	181	WASQVDOGANFSEVSNTSFELNSENVTMKVSVLYNVNTINNTYSCEMIENDIAKATGDIKV	240
Qy	241	TESEIKRSHQLLNKSKSLCVSSFFAISWALLPLSPYMLK	282
Db	241	TESEIKRSHQLLNKSKSLCVSSFFAISWALLPLSPYMLK	282

RESULT 10
US-09-884-441-393
; Sequence 393, Application US/09884441

Patent No. US20020119158A1
GENERAL INFORMATION:
APPLICANT: Algate, Paul A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.462C7
CURRENT APPLICATION NUMBER: 09/067094/884,441
CURRENT FILING DATE: 2001-06-18
NUMBER OF SEQ ID NOS: 489
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 393
LENGTH: 282
TYPE: PRT
ORGANISM: Homo sapiens
US-09-884-441-393

Query Match 100.0%; Score 282; DB 9; Length 282;
Best Local Similarity 100.0%; Pred. No. 2e-263;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASIGQLFWFSIIIIIIAGAIILIGFISGRHSITVTVASAGNIGEDGILSCTFEP 60
Db 1 MASIGQLFWFSIIIIIIAGAIILIGFISGRHSITVTVASAGNIGEDGILSCTFEP 60

Qy 61 DIKLSDIVIOWKEGVLGVHVEFKGKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
Db 61 DIKLSDIVIOWKEGVLGVHVEFKGKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120

Qy 121 QLTDAQTYKCYIIITSGKGNALYKTCGAFSPMEVNDYNASSETLRCEAPRWFPPTVV 180
Db 121 QLTDAQTYKCYIIITSGKGNALYKTCGAFSPMEVNDYNASSETLRCEAPRWFPPTVV 180

Qy 181 WASQVDQANFSEVSNSTSFELSENVTMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
Db 181 WASQVDQANFSEVSNSTSFELSENVTMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240

Qy 241 TESBIKRSHQLLNKASLCVSSFFAISWALLPLSPYMLK 282
Db 241 TESBIKRSHQLLNKASLCVSSFFAISWALLPLSPYMLK 282

RESULT 11
US-09-989-732-291
Sequence 291, Application US/09989732
Patent No. US20020123463A1
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
APPLICANT: Baker, Kevin P.
APPLICANT: Botstein, David
APPLICANT: Desnovers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Fong, Sherman
APPLICANT: Gerber, Hanspeter
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Gurney, Austin L.
APPLICANT: Kljavin, Ivar J.
APPLICANT: Napier, Mary A.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Watanabe, Colin K.
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William I.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same

FILE REFERENCE: P2730P1C57
CURRENT APPLICATION NUMBER: US/09/989,732
CURRENT FILING DATE: 2001-11-19
PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
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PRIOR FILING DATE: 1997-11-12
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;; PRIOR APPLICATION NUMBER: 60/091978
;; PRIOR FILING DATE: 1998-07-07
;; PRIOR APPLICATION NUMBER: 60/091982
;; PRIOR FILING DATE: 1998-07-07
;; PRIOR APPLICATION NUMBER: 60/092182
;; PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 282; DB 9; Length 282;
Best Local Similarity 100.0%; Pred. No. 2e-263;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASLGQILFWSIISIIILAGATLIIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60
Db 1 MASLGQILFWSIISIIILAGATLIIIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60

Qy 61 DIKLSDIVIQWLKEGVLGVHEFKGKDELSEQDEMFRGTAVFADQVIVGNASRLKNV 120
Db 61 DIKLSDIVIQWLKEGVLGVHEFKGKDELSEQDEMFRGTAVFADQVIVGNASRLKNV 120

Qy 121 QLTDTAGTYKCYIITSKKGKGNANLEYKTGAFSMPENVVDYNASSETLCEAPRWFPOQTVV 180
Db 121 QLTDTAGTYKCYIITSKKGKGNANLEYKTGAFSMPENVVDYNASSETLCEAPRWFPOQTVV 180

Qy 181 WASQVDOGANFSEVSNSTSFELNSENVTMKVSVLYNVTINNTYSCMIENDIAKATGDIKV 240
Db 181 WASQVDOGANFSEVSNSTSFELNSENVTMKVSVLYNVTINNTYSCMIENDIAKATGDIKV 240

Qy 241 TESEIKRRSHLQLLNKSKASLCVSSFFAISWALLPLSPYLMK 282
Db 241 TESEIKRRSHLQLLNKSKASLCVSSFFAISWALLPLSPYLMK 282

RESULT 12

US-09-991-073-291
; Sequence 291, Application US/09991073
; Patent No. US20020127576A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gottitsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin L.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Napier, Mary A.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.

APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Watanabe, Colin K.
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William I.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C15
CURRENT APPLICATION NUMBER: US/09/991,073
CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-10-17
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Query Match 100.0%; Score 282; DB 9; Length 282;

Best Local Similarity 100.0%; Pred. No. 2e-263;

Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASIGQILFWSIISIIILAGAIALLIIFGSGRHSITVTTVASAGNIGEDGILSCTFEP 60
 Db 1 MASIGQILFWSIISIIILAGAIALLIIFGSGRHSITVTTVASAGNIGEDGILSCTFEP 60
 Qy 61 DIKLSDIVIOWLKEGVLGLVHEPKEGKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
 Db 61 DIKLSDIVIOWLKEGVLGLVHEPKEGKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
 Qy 121 QLTDAQTYKCVIITSKGNANLEYKGTGAFSMPEVNVVDYNASSETLRCAPRPPQPTVV 180
 Db 121 QLTDAQTYKCVIITSKGNANLEYKGTGAFSMPEVNVVDYNASSETLRCAPRPPQPTVV 180
 Qy 181 WASQVDOGAFSEVSNTSFELSENVTMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
 Db 181 WASQVDOGAFSEVSNTSFELSENVTMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
 Qy 241 TESIERSHQLNLSKASLCVSSFFAISWALLPLSPYMLK 282
 Db 241 TESIERSHQLNLSKASLCVSSFFAISWALLPLSPYMLK 282

RESULT 13

US-09-990-442-291

; Sequence 291, Application US/09990442

; Patent No. US20020132252A1

; GENERAL INFORMATION:

; APPLICANT: Ashkenazi, Avi J.

; APPLICANT: Baker, Kevin P.

; APPLICANT: Botstein, David

; APPLICANT: Deenoyers, Luc

; APPLICANT: Eaton, Dan L.

; APPLICANT: Ferrara, Napoleone

; APPLICANT: Fong, Sherman

; APPLICANT: Gerber, Hanspeter

; APPLICANT: Gerritsen, Mary E.
 ; APPLICANT: Goddard, Audrey
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 ; APPLICANT: Grimaldi, J. Christopher
 ; APPLICANT: Gurney, Austin L.
 ; APPLICANT: Kljavin, Ivar J.
 ; APPLICANT: Napier, Mary A.
 ; APPLICANT: Pan, James
 ; APPLICANT: Paoni, Nicholas F.
 ; APPLICANT: Roy, Margaret Ann
 ; APPLICANT: Stewart, Timothy A.
 ; APPLICANT: Tumas, Daniel
 ; APPLICANT: Watanabe, Colin K.
 ; APPLICANT: Williams, P. Mickey
 ; APPLICANT: Wood, William I.
 ; APPLICANT: Zhang, Zemin
 ; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
 ; FILE REFERENCE: P2730P1C8
 ; CURRENT APPLICATION NUMBER: US/09/990,442
 ; CURRENT FILING DATE: 2001-11-14
 ; PRIOR APPLICATION NUMBER: 60/049787
 ; PRIOR FILING DATE: 1997-06-16
 ; PRIOR APPLICATION NUMBER: 60/062250
 ; PRIOR FILING DATE: 1997-10-17
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;; PRIOR FILING DATE: 1998-07-07
;; PRIOR APPLICATION NUMBER: 60/092182
;; PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 282; DB 9; Length 282;
Best Local Similarity 100.0%; Pred. No. 2e-263;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db 1 MASLGQILFWSIISIIIIILAGATAIIIGFGISGRHSITVTIVASAGNIGEDGILSCTPEP 60
Qy 61 DIKLSDIVIOMLKEGVLGVHFEKGEKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
Db 61 DIKLSDIVIOMLKEGVLGVHFEKGEKDELSEQDEMFRGRTAVFADQVIVGNASRLKNV 120
Qy 121 QLTDAGTYKCYIITSKGGNANLEYKTGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180
Db 121 QLTDAGTYKCYIITSKGGNANLEYKTGAFSMPENVVDYNASSETLRCEAPRFPQPTVV 180
Qy 181 WASQVDOGANFSEVSNVTSFELNSENVTMKVSVLYNTVNTTNTYSCMIENDIAKATGDIKV 240
Db 181 WASQVDOGANFSEVSNVTSFELNSENVTMKVSVLYNTVNTTNTYSCMIENDIAKATGDIKV 240
Qy 241 TSEIKRSHLOLLNSKASLCVSSFFAISWALLPLSPYLMUK 282
Db 241 TSEIKRSHLOLLNSKASLCVSSFFAISWALLPLSPYLMUK 282

RESULT 14
US-09-991-163-291
; Sequence 291, Application US/09991163
; Patent No. US20020132253A1

GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
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; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, J. Christopher
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; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: P2730PIC17
; CURRENT APPLICATION NUMBER: US/09/391,163
; CURRENT FILING DATE: 2001-11-14
; PRIOR APPLICATION NUMBER: 60/045787
; PRIOR FILING DATE: 1997-06-16
; PRIOR APPLICATION NUMBER: 60/062250
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; PRIOR APPLICATION NUMBER: 60/065186
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60/091978	PRIOR APPLICATION NUMBER: 60/091978
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60/091983	PRIOR FILING DATE: 1998-07-07
60/092182	PRIOR APPLICATION NUMBER: 60/092182
60/092183	PRIOR FILING DATE: 1998-07-09

Query Match

Best Local Similarity	100.0%;	Pred. No. 2e-263;	
Matches 282;	Conservative	0;	Mismatches 0;
			Indels 0;
			Gaps 0;

Qy	1	MASLGOILFWISIIIIIIILAGAIALIIIGGISGRHSITVTTVVASAGNIGBGILISCTPEP	60
Db	1	MASLGOILFWISIIIIIIILAGAIALIIIGGISGRHSITVTTVVASAGNIGBGILISCTPEP	60
Qy	61	DIKLSDIVIQWLKEGVIGLVHFEKGGKDBLSQDSMEFRGRTAFVADQOIVGNASIRLKNV	120
Db	61	DIKLSDIVIQWLKEGVIGLVHFEKGGKDBLSQDSMEFRGRTAFVADQOIVGNASIRLKNV	120
Qy	121	QLTDAGTYKCYIIITSKGGKANLEYKTAGFSMPENVVDYNASSETLRCAPRPFQPTTV	180
Db	121	QLTDAGTYKCYIIITSKGGKANLEYKTAGFSMPENVVDYNASSETLRCAPRPFQPTTV	180
Qy	181	WASQVDQGANFSEVSNSTSPELNSENVTKMKVSVLVNVTINNTYSQMIENDIAKATGDIKV	240
Db	181	WASQVDQGANFSEVSNSTSPELNSENVTKMKVSVLVNVTINNTYSQMIENIAKATGDIKV	240

[illegible]

RESULT 15

US-09-993-604-291

Sequence 291, Application US/09993604

Patent No. US20020137075A1

GENERAL INFORMATION:

APPLICANT: Ashkenazi, Avi J.

APPLICANT: Baker, Kevin P.

APPLICANT: Botstein, David

APPLICANT: Desnovers, Luc

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APPLICANT: Watanabe, Colin K.

APPLICANT: Williams, P. Mickey

APPLICANT: Wood, William I.

APPLICANT: Zhang, Zemin

TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic Acids Encoding the Same

FILE REFERENCE: P2730PIC25

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CURRENT FILING DATE: 2001-11-14

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; PRIOR APPLICATION NUMBER: 60/091633
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; PRIOR APPLICATION NUMBER: 60/091978
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match 100.0%; Score 282; DB 9; Length 282;

Best Local Similarity 100.0%; Pred. No. 2e-263;

Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASIGQLFWSIIIIILAGAILIIFGIGSRHSITVTTVASAGNIGEDGILSCTFEP 60

Db 1 MASIGQLFWSIIIIILAGAILIIFGIGSRHSITVTTVASAGNIGEDGILSCTFEP 60

Qy 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEODEMFRGHTAVFADQVIIVGNASLRKNV 120

Db 61 DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEODEMFRGHTAVFADQVIIVGNASLRKNV 120

Qy	121	QLTDAGTYKCYIIITSKGGKGNANLEYKTGAFSMPPEVNVVDYNASSETLRCEAPRWFPQPTVV	180
Db	121	QLTDAGTYKCYIIITSKGGKGNANLEYKTGAFSMPPEVNVVDYNASSETLRCEAPRWFPQPTVV	180
Qy	181	WASQVDOGANFSEVSNTSFELNSENVTKVSVLYNVYIINTYSCMIENDIAKATGDIKV	240
Db	181	WASQVDOGANFSEVSNTSFELNSENVTKVSVLYNVYIINTYSCMIENDIAKATGDIKV	240
Qy	241	TESEIKRSHLQLNSKASLCVSSFFAISWALLPLSPYLMK	282
Db	241	TESEIKRSHLQLNSKASLCVSSFFAISWALLPLSPYLMK	282

Search completed: April 19, 2005, 07:39:45
Job time : 77 secs

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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on: April 19, 2005, 07:17:05 ; Search time 90 Seconds
(without alignments)
1211.849 Million cell updates/sec

Title: us-10-773-715-6

Perfect score: 282

Sequence: 1 MASLGQLFWSIIIIIIIA.....SSFFAISWALLPLSLPYMLK 282

Scoring table: OLIGO

Gapop 60.0 , Gapext 60.0

Searched: 2105692 seqs, 386760381 residues

Word size : 0

Total number of hits satisfying chosen parameters: 2105692

Minimum DB seq length: 0

Maximum DB seq length: 20000000000

Post-processing: Listing first 45 summaries

Database :

- 1: Geneseqp16Dec04.*
- 2: Geneseqp1980s.*
- 3: Geneseqp1990s.*
- 4: Geneseqp2000s.*
- 5: Geneseqp2001s.*
- 6: Geneseqp2002s.*
- 7: Geneseqp2003as.*
- 8: Geneseqp2003bs.*
- 9: Geneseqp2004s.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	282	100.0	282	3	Aay66719 Membrane-
2	282	100.0	282	3	Aab12557 Human ova
3	282	100.0	282	4	Aau29132 Human PRO
4	282	100.0	282	4	Aab87555 Human PRO
5	282	100.0	282	4	Aab99204 Human ova
6	282	100.0	282	4	Aab65242 Human PRO
7	282	100.0	282	5	Aae20311 Human B7-
8	282	100.0	282	5	Abg96445 Human ova
9	282	100.0	282	5	Aau77766 Tumour as
10	282	100.0	282	5	Abg95880 Human sec
11	282	100.0	282	5	Aau76536 Tumour-as
12	282	100.0	282	5	Abp30901 O8E prote
13	282	100.0	282	5	Abb76274 Breast BS
14	282	100.0	282	5	Aae18336 Human B7-
15	282	100.0	282	5	Abb09879 Amino aci
16	282	100.0	282	5	Aae19013 Human B7-
17	282	100.0	282	6	Abu58508 Human PRO
18	282	100.0	282	6	Abu88056 Novel hum
19	282	100.0	282	6	Abu84371 Human sec
20	282	100.0	282	6	Abf66245 Human sec
21	282	100.0	282	6	Abf65635 Human sec
22	282	100.0	282	6	Abu99575 Human sec
23	282	100.0	282	6	Abu58057 Human PRO
24	282	100.0	282	6	Abu59135 Novel hum
25	282	100.0	282	6	Abu82647 Human sec

26	282	100.0	282	6	ABU82814	Human PRO
27	282	100.0	282	6	ABU89935	Novel hum
28	282	100.0	282	6	ABR68184	Human sec
29	282	100.0	282	6	ABU60566	Human sec
30	282	100.0	282	6	ABU96237	Novel hum
31	282	100.0	282	6	ABU92668	Human sec
32	282	100.0	282	6	ABO08745	Human sec
33	282	100.0	282	6	ABO02797	Human sec
34	282	100.0	282	6	ABR74951	Human sec
35	282	100.0	282	6	ABR94713	Human sec
36	282	100.0	282	6	ABU13948	Human PRO
37	282	100.0	282	6	ABU85686	Human PRO
38	282	100.0	282	6	ABU98846	Novel hum
39	282	100.0	282	6	ABU98061	Novel hum
40	282	100.0	282	6	ABU91767	Novel hum
41	282	100.0	282	6	ABU89460	Human PRO
42	282	100.0	282	6	ABU86301	Human sec
43	282	100.0	282	6	ABU67514	Human sec
44	282	100.0	282	6	ABU80542	Human PRO
45	282	100.0	282	6	ABU72533	Novel hum

ALIGNMENTS

RESULT 1

AAy66719

ID AAY66719 standard; protein; 282 AA.

AC AAY66719;

DT 05-APR-2000 (first entry)

DE Membrane-bound protein PRO1291.

XX Membrane-bound polypeptide; PRO polypeptide; LDL receptor; TIE ligand;
pharmaceutical; receptor immunoadhesin; gene mapping.

OS Homo sapiens.

PN WO9963088-A2.

PD 09-DEC-1999.

PF 02-JUN-1999; 99WO-US012252.

PR 02-JUN-1998; 98US-0087607P.

PR 02-JUN-1998; 98US-0087759P.

PR 03-JUN-1998; 98US-0087827P.

PR 04-JUN-1998; 98US-0088021P.

PR 04-JUN-1998; 98US-0088025P.

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PR 04-JUN-1998; 98US-0088029P.

PR 04-JUN-1998; 98US-0088030P.

PR 04-JUN-1998; 98US-0088033P.

PR 05-JUN-1998; 98US-0088167P.

PR 05-JUN-1998; 98US-0088202P.

PR 05-JUN-1998; 98US-0088212P.

PR 05-JUN-1998; 98US-0088217P.

PR 09-JUN-1998; 98US-0088655P.

PR 10-JUN-1998; 98US-0088722P.

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PR 10-JUN-1998; 98US-0088741P.

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 PR 07-JUL-1998; 98US-0091978P.
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 PR 10-JUL-1998; 98US-0092182P.
 PR 10-JUL-1998; 98US-0092472P.
 PR 20-JUL-1998; 98US-0093339P.
 PR 30-JUL-1998; 98US-0094511P.
 PR 04-AUG-1998; 98US-0095282P.
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 PR 04-AUG-1998; 98US-0095301P.
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 PR 10-AUG-1998; 98US-00953916P.

PR 10-AUG-1998; 98US-0095929P.
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 PR 26-AUG-1998; 98US-0098014P.
 PR 31-AUG-1998; 98US-0098525P.
 PR 16-SEP-1998; 98US-0100634P.
 PR 12-JAN-1999; 99US-0115565P.
 XX
 (GETH) GENENTECH INC.

Baker K, Chen J, Goddard A, Gurney AL, Smith V, Watanabe CK;
 Wood WI, Yuan J;

WPI; 2000-072883/06.
 N-PSDB; AAZ65059.

Membrane-bound proteins and related nucleotide sequences.

Claim 12; Fig 208; 822pp; English.

CC The invention provides membrane-bound PRO polypeptides and polynucleotides encoding them. The PRO sequences of the invention were identified based on extracellular domain homology screening. The PRO sequences have homology with proteins including LDL receptors, TIE ligands and various enzymes. The membrane-bound proteins and receptor molecules are useful as pharmaceutical and diagnostic agents. Receptor immunoadhesins, for instance, can be used as therapeutic agents to block receptor-ligand interactions. The membrane-bound proteins can also be employed for screening of potential peptide or small molecule inhibitors of the relevant receptor/ligand interaction. The PRO encoding sequences are useful as hybridization probes, in chromosome and gene mapping and in the generation of antisense RNA and DNA. PRO nucleic acid sequences will also be useful for the preparation of PRO polypeptides, especially by recombinant techniques

SQ Sequence 282 AA;

Query Match 100.0%; Score 282; DB 3; Length 282;
 Best Local Similarity 100.0%; Pred. No. 2,8e-286;

Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MASLQILFWSIISIIIIILAGAIILIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60

DB 1 MASLQILFWSIISIIIIILAGAIILIGFISGRHSITVTTVASAGNIGEDGILSCTFEP 60

QY 61 DIKLSDIVIOWLKEGVGLVHEFKGKDELSEODEMFRGRTAVFADQVIVGNASRLKNV 120
Db 61 DIKLSDIVIOWLKEGVGLVHEFKGKDELSEODEMFRGRTAVFADQVIVGNASRLKNV 120
QY 121 QLTDAAGTYKCYIITSKKGKGNANLEYKTGAFSPMEPVNDYNNASSETLCEAPRWFPQPTVV 180
Db 121 QLTDAAGTYKCYIITSKKGKGNANLEYKTGAFSPMEPVNDYNNASSETLCEAPRWFPQPTVV 180
QY 181 WASQVDOGANFSEVSNSTSPFELNSENVMTKVVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
Db 181 WASQVDOGANFSEVSNSTSPFELNSENVMTKVVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
QY 241 TESEIKRRSHQLLNSKASLCVSSFFAISWALLPLSPYLMLK 282
Db 241 TESEIKRRSHQLLNSKASLCVSSFFAISWALLPLSPYLMLK 282
RESULT 2
AAB12557
ID AAB12557 standard; protein; 282 AA.
XX
AC AAB12557;
XX
DT 07-NOV-2000 (first entry)
XX
DE Human ovarian carcinoma antigen O8E protein SEQ ID NO:393.
XX
KW Human; ovarian carcinoma; ovarian cancer; therapy; diagnosis;
KW tumour antigen; identification; cytostatic; gene therapy; vaccine.
XX
OS Homo sapiens.
XX
PN WO200036107-A2.
XX
PD 22-JUN-2000.
XX
PF 17-DEC-1999; 99WO-US030270.
XX
PR 17-DEC-1998; 98US-00215681.
PR 17-DEC-1998; 98US-00216003.
PR 23-JUN-1999; 99US-00338933.
PR 24-SEP-1999; 99US-00404679.
XX
PA (CORI-) CORIXA CORP.
XX
PI Mitcham JL, King GE, Algate PA, Frudakis TN;
XX
XX
DR WPI; 2000-431589/37.
XX
PT Immunogenic portion of an ovarian carcinoma protein and the nucleic acid
PT encoding it, useful for the diagnosis, prevention and treatment of
PT cancer, preferably ovarian cancer.
XX
PS Example 2; Page 207; 299pp; English.
XX
CC The present invention describes an isolated polypeptide comprising an
CC immunogenic portion of an ovarian carcinoma protein (or its variants).
CC Ovarian carcinoma proteins, and polynucleotides encoding them, have
CC cytotatic activity and can be used in gene therapy and vaccines. Ovarian
CC carcinoma polypeptides, nucleic acids, antibodies and vaccines are useful
CC for the prevention, diagnosis and treatment of cancer, preferably ovarian
CC cancer. AAG9691 to AAA70077 and AAB12552 to AAB12557 represent human
CC ovarian carcinoma polynucleotides and proteins used in the
CC exemplification of the present invention
XX
SQ Sequence 282 AA;
Query Match 100.0%; Score 282; DB 3; Length 282;
Best Local Similarity 100.0%; Pred. No. 2.8e-286; Indels 0; Gaps 0;
Matches 282; Conservative 0; Mismatches 0;
QY 1 MASLGQILFWSIISIIILAGAIALIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP 60

Db 1 MASLGQILFWSIISIIILAGAIALIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP 60
QY 61 DIKLSDIVIOWLKEGVGLVHEFKGKDELSEODEMFRGRTAVFADQVIVGNASRLKNV 120
Db 61 DIKLSDIVIOWLKEGVGLVHEFKGKDELSEODEMFRGRTAVFADQVIVGNASRLKNV 120
QY 121 QLTDAAGTYKCYIITSKKGKGNANLEYKTGAFSPMEPVNDYNNASSETLCEAPRWFPQPTVV 180
Db 121 QLTDAAGTYKCYIITSKKGKGNANLEYKTGAFSPMEPVNDYNNASSETLCEAPRWFPQPTVV 180
QY 181 WASQVDOGANFSEVSNSTSPFELNSENVMTKVVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
Db 181 WASQVDOGANFSEVSNSTSPFELNSENVMTKVVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
QY 241 TESEIKRRSHQLLNSKASLCVSSFFAISWALLPLSPYLMLK 282
Db 241 TESEIKRRSHQLLNSKASLCVSSFFAISWALLPLSPYLMLK 282
RESULT 3
AAU29132
ID AAU29132 standard; protein; 282 AA.
XX
AC AAU29132;
XX
DT 18-DEC-2001. (first entry)
XX
DE Human PRO polypeptide sequence #109.
XX
KW PRO polypeptide; mammal; tumour; cancer; human; cattle; horse; sheep;
KW dog; cat; pig; goat; rabbit; tumour necrosis factor alpha; TNF-alpha;
KW blood; chondrocyte cell; cell proliferation; cell differentiation; colon;
KW adrenal; lung; breast; prostate; rectum; cervix; liver; genetic disorder.
XX
OS Homo sapiens.
XX
PN WO200168848-A2.
XX
PD 20-SEP-2001.
XX
PF 28-FEB-2001; 2001WO-US006520.
XX
PR 01-MAR-2000; 2000WO-US005601.
PR 02-MAR-2000; 2000WO-US005841.
PR 03-MAR-2000; 2000US-0187202P.
PR 06-MAR-2000; 2000US-0186968P.
PR 14-MAR-2000; 2000US-0189320P.
PR 14-MAR-2000; 2000US-0189328P.
PR 15-MAR-2000; 2000WO-US006884.
PR 21-MAR-2000; 2000US-0190828P.
PR 21-MAR-2000; 2000US-0191007P.
PR 21-MAR-2000; 2000US-0191048P.
PR 21-MAR-2000; 2000US-0191314P.
PR 28-MAR-2000; 2000US-0192855P.
PR 29-MAR-2000; 2000US-0193032P.
PR 30-MAR-2000; 2000US-0193053P.
PR 30-MAR-2000; 2000WO-US008439.
PR 04-APR-2000; 2000US-0194449P.
PR 04-APR-2000; 2000US-0194647P.
PR 11-APR-2000; 2000US-0195975P.
PR 11-APR-2000; 2000US-0196000P.
PR 11-APR-2000; 2000US-0196187P.
PR 11-APR-2000; 2000US-0196690P.
PR 11-APR-2000; 2000US-0196820P.
PR 18-APR-2000; 2000US-0198121P.
PR 18-APR-2000; 2000US-0198585P.
PR 25-APR-2000; 2000US-0199397P.
PR 25-APR-2000; 2000US-0199550P.
PR 25-APR-2000; 2000US-0199654P.
PR 03-MAY-2000; 2000US-0201516P.
PR 17-MAY-2000; 2000WO-US013705.
PR 22-MAY-2000; 2000WO-US014042.

PR 30-MAY-2000; 2000WO-US014941.
 PR 02-JUN-2000; 2000WO-US015264.
 PR 05-JUN-2000; 2000US-0209832P.
 PR 28-JUL-2000; 2000WO-US020710.
 PR 22-AUG-2000; 2000US-00644848.
 PR 24-AUG-2000; 2000WO-US023328.
 PR 08-NOV-2000; 2000WO-US030952.
 PR 01-DEC-2000; 2000WO-US032678.
 PR 20-DEC-2000; 2000WO-US034956.
 XX
 PA (GETH) GENENTECH INC.
 XX
 XX Baker KP, Chen J, Deanoyers L, Goddard A, Godowski PJ, Gurney AL;
 PI Pan J, Smith V, Watanabe CK, Wood WI, Zhang Z;
 XX
 DR WPI; 2001-602746/68.
 DR N-PSDB; AAS46033.
 XX
 PT Novel nucleic acids encoding PRO polypeptides, used to diagnose the
 PT presence of tumors, such as prostate and breast tumors, in mammals and to
 PT screen for modulators of the compounds.
 XX
 PS Claim 11; Fig 218; 774pp; English.
 XX
 CC Sequences AAU29024-AAU29328 represent PRO polypeptides of the invention.
 CC The PRO polypeptides and their associated nucleic acids can be used to
 CC detect the presence of a tumour in a mammal by comparing the level of
 CC expression of a PRO polypeptide in a test sample of cells from the animal
 CC and a control sample of normal cells, whereby a higher level of
 CC expression in the test sample indicates the presence of a tumour in the
 CC mammal. Mammals include dogs, cats, cattle, horses, sheep, pigs, goats
 CC and rabbits but are preferably human. The polypeptides can be used to
 CC stimulate tumour necrosis factor (TNF) alpha release from human blood,
 CC when contacted with it. A specific polypeptide can be used to stimulate
 CC the proliferation or differentiation of chondrocyte cells. The PRO
 CC proteins can be used to determine the presence of tumours and also
 CC susceptibility to tumour development, particularly adrenal, lung, colon,
 CC breast, prostate, cervical, or liver tumours, in mammalian
 CC subjects. The oligonucleotide probes specific for the PRO nucleic acids
 CC can be used for genetic analysis of individuals with genetic disorders
 XX
 SQ Sequence 282 AA;
 PS
 Query Match 100.0%; Score 282; DB 4; Length 282;
 Best Local Similarity 100.0%; Pred. No. 2.8e-286;
 Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 MASLGQLFWSIIISIIILAGAILIIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP 60
 DB 1 MASLGQLFWSIIISIIILAGAILIIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP 60
 QY 61 DIKLSDIVIOWLKEGVLGVLHFEKGEKDELSEQDEMFRGRTAVFADQVIVGNASLRLKNV 120
 DB 61 DIKLSDIVIOWLKEGVLGVLHFEKGEKDELSEQDEMFRGRTAVFADQVIVGNASLRLKNV 120
 QY 121 QLTDAAGTYKCYIITSKGNANLEYKTGAFSPMEVNVNDYNASSETLRCEAPRFPPTVV 180
 DB 121 QLTDAAGTYKCYIITSKGNANLEYKTGAFSPMEVNVNDYNASSETLRCEAPRFPPTVV 180
 QY 181 WASQVDQGANFSEVSNSTFELNSENVTKVSVLVNNTYSCMIENDIAKATGDIKV 240
 DB 181 WASQVDQGANFSEVSNSTFELNSENVTKVSVLVNNTYSCMIENDIAKATGDIKV 240
 QY 241 TESEIKRRSHQLNLNSKASLCVSSFFAISWALLPLSPYMLK 282
 DB 241 TESEIKRRSHQLNLNSKASLCVSSFFAISWALLPLSPYMLK 282
 RESULT 4
 AAB87555
 ID AAB87555 standard; protein; 282 AA.
 XX
 AC AAB87555;

XX 15-MAY-2001 (first entry)
 XX Human PRO1291.
 DE
 XX Human, PRO protein; mapping.
 KW
 XX Homo sapiens.
 OS
 XX WO200116318-A2.
 PN
 XX 08-MAR-2001.
 PD
 XX 24-AUG-2000; 2000WO-US023328.
 PF
 XX 01-SEP-1999; 99WO-US020111.
 PR 15-SEP-1999; 99WO-US021090.
 PR 07-DEC-1999; 99US-0169495P.
 PR 09-DEC-1999; 99US-0170262P.
 PR 11-JAN-2000; 2000US-0175481P.
 PR 18-FEB-2000; 2000WO-US004341.
 PR 22-FEB-2000; 2000WO-US004414.
 PR 01-MAR-2000; 2000WO-US005601.
 PR 03-MAR-2000; 2000US-0187202P.
 PR 21-MAR-2000; 2000US-0191007P.
 PR 30-MAR-2000; 2000WO-US008439.
 PR 25-APR-2000; 2000US-0199397P.
 PR 22-MAY-2000; 2000WO-US014042.
 PR 05-JUN-2000; 2000US-0209832P.
 XX
 PA (GETH) GENENTECH INC.
 XX
 XX Eaton DL, Filvaroff E, Gerritsen ME, Goddard A, Godowski PJ;
 PI Grimaldi CJ, Gurney AL, Watanabe CK, Wood WI;
 XX
 DR WPI; 2001-183260/18.
 DR N-PSDB; AAF92087.
 XX
 PT Eighty four nucleic acids encoding PRO polypeptides, useful in molecular
 PT biology, including use as hybridization probes, and in chromosome and
 PT gene mapping.
 XX
 PS Claim 12; Fig 60; 278pp; English.
 XX
 CC The present sequence is a human PRO polypeptide (secreted and
 CC transmembrane). The PRO protein, and PRO agonists, PRO antagonists or
 CC anti-PRO antibodies are useful for preparation of a medicament useful in
 CC the treatment of a condition which is responsive to the PRO protein,
 CC agonists, antagonists or anti-PRO antibodies. The PRO protein may also be
 CC employed as molecular weight markers for protein electrophoresis. The PRO
 CC coding sequence has applications in molecular biology, including use as
 CC hybridisation probes, and in chromosome and gene mapping
 XX
 SQ Sequence 282 AA;
 PS
 Query Match 100.0%; Score 282; DB 4; Length 282;
 Best Local Similarity 100.0%; Pred. No. 2.8e-286;
 Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 MASLGQLFWSIIISIIILAGAILIIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP 60
 DB 1 MASLGQLFWSIIISIIILAGAILIIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP 60
 QY 61 DIKLSDIVIOWLKEGVLGVLHFEKGEKDELSEQDEMFRGRTAVFADQVIVGNASLRLKNV 120
 DB 61 DIKLSDIVIOWLKEGVLGVLHFEKGEKDELSEQDEMFRGRTAVFADQVIVGNASLRLKNV 120
 QY 121 QLTDAAGTYKCYIITSKGNANLEYKTGAFSPMEVNVNDYNASSETLRCEAPRFPPTVV 180
 DB 121 QLTDAAGTYKCYIITSKGNANLEYKTGAFSPMEVNVNDYNASSETLRCEAPRFPPTVV 180
 QY 181 WASQVDQGANFSEVSNSTFELNSENVTKVSVLVNNTYSCMIENDIAKATGDIKV 240

```
Db 181 WASQVDOGANFSEVSNSTFELNSENVTKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
    |||||||
Qy 241 TESIERSHLQLLNKSKSLCVSFFFAISWALLPLSPYMLK 282
    |||||||
Db 241 TESIERSHLQLLNKSKSLCVSFFFAISWALLPLSPYMLK 282
    |||||||

RESULT 5
AAB99204
ID AAB99204 standard; protein; 282 AA.
AC AAB99204;
XX
DT 04-SEP-2001 (first entry)
DE Human ovarian tumour-derived antigen O8E #1.
XX
KW Cytostatic; human; breast tumour protein; breast cancer; ovarian tumour;
KW antigen; O8E.
XX
OS Homo sapiens.
XX
PN WO200140269-A2.
XX
PD 07-JUN-2001.
XX
PF 29-NOV-2000; 2000WO-US032520.
XX
PR 30-NOV-1999; 99US-00451651.
PR 22-FEB-2000; 2000US-00510862.
PR 10-MAR-2000; 2000US-00523586.
PR 07-APR-2000; 2000US-00545068.
PR 15-MAY-2000; 2000US-00571025.
XX
PA (CORI-) CORIXA CORP.
XX
PI Dillon DC, Day CH, Jiang Y, Houghton RL, Mitcham JL, Wang A;
XX
DR WPI; 2001-356154/37.
XX
DR N-PSDB; AAH55681.
XX
PT Breast tumor polypeptides and the nucleic acids that encode them, useful
PT for the prevention, diagnosis and treatment of breast cancer.
XX
PS Example 3; Page 190; 221pp; English.
XX
CC The present invention relates to human breast tumour protein coding
CC sequences (see AAH55479-AAH55513, AAH55517-AAH55679 and AAH55682-
CC AAH55762). The breast tumour protein DNA sequences may be used in the
CC prevention, diagnosis and treatment of diseases associated with
CC inappropriate expression of the breast tumour protein e.g. breast cancer.
CC The present sequence is a human ovarian tumour-derived antigen, which was
CC used in an example from the present invention
XX
SQ Sequence 282 AA;

Query Match 100.0%; Score 282; DB 4; Length 282;
Best Local Similarity 100.0%; Pred. No. 2.8e-286;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASLGQILFWSIIISIIILAGAIALLIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP 60
    |||||||
Db 1 MASLGQILFWSIIISIIILAGAIALLIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP 60
    |||||||

Qy 61 DIKLSDIVIQWKEGVLGVHFEKKGDELSEQDEMFRGRTAVFADQVIVGNASLRKNV 120
    |||||||
Db 61 DIKLSDIVIQWKEGVLGVHFEKKGDELSEQDEMFRGRTAVFADQVIVGNASLRKNV 120
    |||||||

Qy 121 QLTDAQTYKCVIITSKGGKANLKYKTCAFSPMPENVVDYNASSETLRCEAPRWFPQPTVV 180
    |||||||
Db 121 QLTDAQTYKCVIITSKGGKANLKYKTCAFSPMPENVVDYNASSETLRCEAPRWFPQPTVV 180
    |||||||
```

```
Qy 181 WASQVDOGANFSEVSNSTFELNSENVTKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
    |||||||
Db 181 WASQVDOGANFSEVSNSTFELNSENVTKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
    |||||||

Qy 241 TESIERSHLQLLNKSKSLCVSFFFAISWALLPLSPYMLK 282
    |||||||
Db 241 TESIERSHLQLLNKSKSLCVSFFFAISWALLPLSPYMLK 282
    |||||||

RESULT 6
AAB65242
ID AAB65242 standard; protein; 282 AA.
XX
AC AAB65242;
XX
DT 02-APR-2001 (first entry)
DE Human PR01291 (UNQ659) protein sequence SEQ ID NO:291.
XX
KW Human; secreted and transmembrane protein; PRO; cytostatic; cell death;
KW cancer; chromosomal mapping; gene mapping; tissue typing;
KW diagnostic assay.
XX
OS Homo sapiens.
XX
PN WO2000073454-A1.
XX
PD 07-DEC-2000.
XX
PF 30-MAR-2000; 2000WO-US008439.
XX
PR 02-JUN-1999; 99WO-US012252.
PR 23-JUN-1999; 99US-0141037P.
PR 07-JUL-1999; 99US-0143048P.
PR 20-JUL-1999; 99US-0144758P.
PR 26-JUL-1999; 99US-0145698P.
PR 28-JUL-1999; 99US-0146222P.
PR 17-AUG-1999; 99US-0149396P.
PR 15-SEP-1999; 99WO-US021090.
PR 08-OCT-1999; 99US-0158663P.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 05-JAN-2000; 2000WO-US000219.
PR 06-JAN-2000; 2000WO-US000376.
PR 11-FEB-2000; 2000WO-US003565.
PR 18-FEB-2000; 2000WO-US004341.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US004914.
PR 02-FEB-2000; 2000WO-US005004.
PR 15-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US006884.
PR 20-MAR-2000; 2000WO-US007377.
XX
PA (GETH ) GENENTECH INC.
XX
Qy Ashkenazi AJ, Baker KP, Borstein D, Desnoyers L, Eaton DL;
Qy Ferrara N, Fong S, Gerber H, Gerritsen ME, Goddard A, Godowski PJ;
Qy Girmaldi CJ, Gurney AL, Kljavin IJ, Napier MA, Pan J, Paoni NF;
Qy Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM, Wood WJ;
Qy Zhang Z;
XX
DR WPI; 2001-032160/04.
DR N-PSDB; AAP44205.
XX
PT PRO polynucleotides used to produce polypeptides used to target bioactive
PT molecules such as toxins, radiolabels or antibodies, to specific cells,
PT to cause targeted cell death.
XX
PS Claim 12; Fig 208; 935pp; English.
XX
```

CC The present invention describes human secreted and transmembrane PRO
 CC proteins. The PRO proteins have cytotostatic activity. The PRO proteins can
 CC be used for targeted delivery of bioactive molecules, such as toxins,
 CC radiolabels or antibodies, that cause cell death. PRO nucleotide
 CC sequences, and their fragments, can be used as hybridisation probes, in
 CC chromosomal and gene mapping, and in the generation of anti-sense RNA and
 CC DNA. They may also be used to produce transgenic animals which are used
 CC to develop and screen therapeutically useful reagents. The PRO nucleotide
 CC and protein sequence can be used for tissue typing and in treating
 CC cancer. Anti-PRO antibodies can be used in diagnostic assays. AAF44270 to
 CC AAF44470 represent PCR primers and hybridisation probes used in the
 CC isolation of human PRO sequences. AAF44087 to AAF44269 and AAB65154 to
 CC AAB65300 represent human PRO polynucleotide and protein sequences given
 CC in the exemplification of the present invention
 XX
 SQ Sequence 282 AA;

Query Match 100.0%; Score 282; DB 4; Length 282;
 Best Local Similarity 100.0%; Pred. No. 2.8e-286;
 Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASLGQLFWISIIIIILAGATIALIIGFGISGRHSITVTVASAGNIGDGLSCTFEP 60
 Db |||||
 Qy 1 MASLGQLFWISIIIIILAGATIALIIGFGISGRHSITVTVASAGNIGDGLSCTFEP 60
 Db |||||
 Qy 61 DIKLSDIVIOWLKEGVLGLVHEFKEGKDELSEODEMFRGRTAVPADQVIVGNASLRKNV 120
 Db |||||
 Qy 61 DIKLSDIVIOWLKEGVLGLVHEFKEGKDELSEODEMFRGRTAVPADQVIVGNASLRKNV 120
 Db |||||
 Qy 121 QLTDAAGTYKCYIITSKGNANLEYKTFGAFSMPVNVVDYNASSETLRCEAPRFPQPTVV 180
 Db |||||
 Qy 121 QLTDAAGTYKCYIITSKGNANLEYKTFGAFSMPVNVVDYNASSETLRCEAPRFPQPTVV 180
 Db |||||
 Qy 181 WASQVDOGANFSEVSNVTSFELNSNTVMKVSVLYNVYNTTSCMIENDIAKATGDIKV 240
 Db |||||
 Qy 181 WASQVDOGANFSEVSNVTSFELNSNTVMKVSVLYNVYNTTSCMIENDIAKATGDIKV 240
 Db |||||
 Qy 241 TESEIKRRSHLOLLNSKASLCVSSFFAISWALLPLSPYLMK 282
 Db |||||
 Qy 241 TESEIKRRSHLOLLNSKASLCVSSFFAISWALLPLSPYLMK 282
 Db |||||

RESULT 7
 AAE20311
 ID AAE20311 standard; protein; 282 AA.

XX AAE20311;

XX 18-JUN-2002 (first entry)

XX Human B7-H8 protein #1.

XX Human; B7-like protein; inflammation; tissue damage; immune disorder;
 KW Addison's disease; autoimmune haemolytic anaemia; autoimmune thyroiditis;
 KW diabetes mellitus; Crohn's disease; multiple sclerosis; allergy; cancer;
 KW rheumatoid arthritis; cardiovascular disorder; nervous system disorder;
 KW myocardial ischaemia; ulcerative colitis; reproductive system disorder;
 KW Alzheimer's disease; Parkinson's disease; endocrine disorder; hepatitis;
 KW diabetes mellitus; Grave's disease; Paget's disease; liver disorder;
 KW gastrointestinal disorder; irritable bowel syndrome; cerebral anoxia;
 KW dysphagia; hepatomegaly; neurological disease; infectious disease;
 KW epilepsy; gene therapy; B7-H8 protein; chromosome 1.

OS Homo sapiens.

XX Key Location/Qualifiers
 FH Peptide 1..24
 FT FT
 FT Protein 25..282
 FT /note= "Mature B7-H8 protein"

XX WO200202587-A1.

XX

PD 10-JAN-2002.

XX 29-JUN-2001; 2001WO-US020917.

XX 30-JUN-2000; 2000US-0215135P.

PR 14-AUG-2000; 2000US-0225266P.

XX (HUMA-) HUMAN GENOME SCI INC.

XX Fiscella M, Ni J, Ruben SM;

XX WPI; 2002-257198/30.

DR N-PSDB; AAD32519.

XX Isolated nucleic acids encoding human B7-like polypeptides, useful for
 PT diagnosis and treatment of e.g. inflammation, cancer, immune disorders
 PT such as Addison's disease, and cardiovascular disorders such as
 PT myocardial ischemias.

XX Example 1; Fig 1; 493pp; English.

XX The present invention relates to novel human B7-like polypeptides and
 CC polynucleotides encoding such proteins. Sequences of the invention are
 CC used for preventing, treating or ameliorating a medical condition in a
 CC mammalian subject. The polynucleotides and polypeptides are administered
 CC to subjects having a disorder related to B-7 Like polypeptides, such as
 CC inappropriate or excessive inflammation which can lead to tissue damage
 CC or even death, where the inflammation is brought about by the activation
 CC of certain cells in the body e.g. T cells and may involve disorders
 CC related to immune system. The nucleic acids, proteins, antibodies,
 CC agonists and antagonists of the invention are useful in the diagnosis,
 CC treatment and prevention of cancer (e.g. cancers of the adrenal gland,
 CC bone, bone marrow, breast, gastrointestinal tract, liver, urogenital or
 CC lung), immune disorders (e.g., Addison's disease, allergies, autoimmune
 CC haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's
 CC disease, multiple sclerosis, rheumatoid arthritis, ulcerative colitis,
 CC cardiovascular disorders (e.g., myocardial ischaemia), nervous system
 CC disorders (Alzheimer's disease, Parkinson's disease), endocrine disorders
 CC (e.g., diabetes mellitus, Grave's disease), reproductive system disorders
 CC (e.g., cryptorchism, Paget's disease), gastrointestinal disorders (e.g.,
 CC dysphagia, irritable bowel syndrome), liver disorders (e.g., hepatitis,
 CC hepatomegaly), neurological diseases (e.g., cerebral anoxia and epilepsy)
 CC and infectious diseases such as viral, bacterial, fungal and parasitic
 CC infections. Sequences of the invention are also used in gene therapy. The
 CC present sequence is human B7-H8 protein. B7-H8 gene is located on
 CC chromosome 1

XX Sequence 282 AA;

Query Match 100.0%; Score 282; DB 5; Length 282;
 Best Local Similarity 100.0%; Pred. No. 2.8e-286;
 Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASLGQLFWISIIIIILAGATIALIIGFGISGRHSITVTVASAGNIGDGLSCTFEP 60

Db 1 MASLGQLFWISIIIIILAGATIALIIGFGISGRHSITVTVASAGNIGDGLSCTFEP 60

Qy 61 DIKLSDIVIOWLKEGVLGLVHEFKEGKDELSEODEMFRGRTAVPADQVIVGNASLRKNV 120

Db 61 DIKLSDIVIOWLKEGVLGLVHEFKEGKDELSEODEMFRGRTAVPADQVIVGNASLRKNV 120

Qy 121 QLTDAAGTYKCYIITSKGNANLEYKTFGAFSMPVNVVDYNASSETLRCEAPRFPQPTVV 180

Db 121 QLTDAAGTYKCYIITSKGNANLEYKTFGAFSMPVNVVDYNASSETLRCEAPRFPQPTVV 180

Qy 181 WASQVDOGANFSEVSNVTSFELNSNTVMKVSVLYNVYNTTSCMIENDIAKATGDIKV 240

Db 181 WASQVDOGANFSEVSNVTSFELNSNTVMKVSVLYNVYNTTSCMIENDIAKATGDIKV 240

Qy 241 TESEIKRRSHLOLLNSKASLCVSSFFAISWALLPLSPYLMK 282

Db 241 TESEIKRRSHLOLLNSKASLCVSSFFAISWALLPLSPYLMK 282

```
RESULT 8
ABG96445
ID   ABG96445 standard; protein; 282 AA.
XX
XX
XX   ABG96445;
AC
XX
DT   11-DEC-2002 (first entry)
XX
XX   Human ovarian cancer marker OV88.
DE
XX
XX   Human; ovarian cancer; marker; familial history; brain disorder;
XX   central nervous system disorder; bacterial meningitis; viral meningitis;
XX   Alzheimer's disease; Parkinson's disease; cerebral edema; hydrocephalus;
XX   brain herniation; inflammation; encephalitis; testicular disorder;
XX   nontuberculous granulomatous orchitis; connective tissue disorder;
XX   heart disorder; ischaemic heart disease; atherosclerosis; neoplasm;
XX   histological type; carcinogenic; ovarian cancer marker.
XX
OS   Homo sapiens.
XX
XX   WO200271928-A2.
XX
XX   19-SEP-2002.
XX
XX   14-MAR-2002; 2002WO-US007826.
XX
XX   14-MAR-2001; 2001US-0276025P.
XX
XX   14-MAR-2001; 2001US-0276026P.
XX
XX   10-AUG-2001; 2001US-0311732P.
XX
XX   19-SEP-2001; 2001US-0323580P.
XX
XX   26-SEP-2001; 2001US-0324967P.
XX
XX   26-SEP-2001; 2001US-0325102P.
XX
XX   26-SEP-2001; 2001US-0325149P.
XX
XX   (MILL-) MILLENNIUM PHARM INC.
XX
XX   Monahan JE, Gannavarapu M, Hoersch S, Kamatkar S, Kovatis SG;
XX   Meyers RE, Morrissey MP, Olandt PJ, Sen A, Vieby PO, Mills GB;
XX   Bast RC, Lu K, Schmandt RE, Zhao X, Glatt K;
XX
XX   WPI; 2002-723277/78.
XX
XX   N-PSDB; ABS76544.
XX
XX   Assessing whether a patient is afflicted with ovarian cancer, useful in
XX   assessing the stage or progression of the disease, comprises comparing
XX   the expression level of a cancer marker in a sample from a patient and
XX   from a non cancer patient.
XX
XX   Disclosure; Page 468-469; 481pp; English.
XX
XX   The present invention relates to a new method for assessing whether a
XX   patient is afflicted with ovarian cancer. The method involves comparing
XX   the expression level of a marker in a patient sample and the normal level
XX   of expression of the marker in a control non-ovarian cancer sample, where
XX   the marker is selected from 363 cancer markers described in the
XX   specification. The method of the invention is useful in diagnosing or
XX   characterising cancer, in detecting the presence of cancer as early as
XX   possible, and the recurrence of ovarian cancer. The method may also be of
XX   particular use with patients having an enhanced risk of developing
XX   ovarian cancer (e.g. patients having a familial history of ovarian
XX   cancer). The cancer markers may be used in the management and treatment
XX   of e.g. brain and central nervous system disorders (e.g. bacterial and
XX   viral meningitis, Alzheimer's disease or Parkinson's disease), brain
XX   disorders (e.g. cerebral edema, hydrocephalus or brain herniations),
XX   inflammations (e.g. bacterial or viral meningitis or encephalitis),
XX   testicular disorders (e.g. nontuberculous granulomatous orchitis),
XX   connective tissue disorders, or heart disorders (e.g. ischaemic heart
XX   disease or atherosclerosis). The compositions and methods may also be
XX   used in assessing the histological type of neoplasm associated with
XX   ovarian cancer, monitoring the progression of ovarian cancer, determining
XX   whether ovarian cancer has metastasized or is likely to metastasize,
XX   selecting a composition for inhibiting ovarian cancer, assessing the
```

```
CC   ovarian carcinogenic potential of a compound, or inhibiting ovarian
CC   cancer or at risk of developing ovarian cancer. The present amino acid
CC   sequence represents one of the ovarian cancer markers described in the
CC   invention
XX
XX   SQ   Sequence 282 AA;
XX
XX   Query Match      100.0%; Score 282; DB 5; Length 282;
XX   Best Local Similarity 100.0%; Pred. NO. 2.8e-286;
XX   Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
XX
XX   Qy   1 MASIGQLFWSIISIIILAGATALLIGFGISGRHSITVTTVASAGNIGEDGILSCFEP 60
XX   Db   1 MASIGQLFWSIISIIILAGATALLIGFGISGRHSITVTTVASAGNIGEDGILSCFEP 60
XX
XX   Qy   61 DIKLSDIVIOWLKEGVILGLVHEFKEGKDELSEODEMFRGRTAVFADQVIVGNASLRLKNV 120
XX   Db   61 DIKLSDIVIOWLKEGVILGLVHEFKEGKDELSEODEMFRGRTAVFADQVIVGNASLRLKNV 120
XX
XX   Qy   121 QLTDAGTYKCYIITSKGGNANLEYKTGAFSMPEVNVVDYNASSETLRCEAPRWFPQPTVV 180
XX   Db   121 QLTDAGTYKCYIITSKGGNANLEYKTGAFSMPEVNVVDYNASSETLRCEAPRWFPQPTVV 180
XX
XX   Qy   181 WASQVDOGANFSEVSNTSFELNSENVTMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
XX   Db   181 WASQVDOGANFSEVSNTSFELNSENVTMKVSVLYNVNTINNTYSCMIENDIAKATGDIKV 240
XX
XX   Qy   241 TESEIKRSHQLLNKSKASLCVSVFFAISWALLPLSPYLMK 282
XX   Db   241 TESEIKRSHQLLNKSKASLCVSVFFAISWALLPLSPYLMK 282
XX
XX   RESULT 9
XX   AAU77766
XX   ID   AAU77766 standard; protein; 282 AA.
XX
XX   AC   AAU77766;
XX
XX   DT   05-JUN-2002 (first entry)
XX
XX   DE   Tumour associated antigenic target polypeptide (TAT) 136.
XX
XX   KW   Tumour associated antigenic target polypeptide; TAT; cancer;
XX   breast cancer; colorectal cancer; lung cancer; ovarian cancer;
XX   central nervous system cancer; liver cancer; bladder cancer;
XX   pancreatic cancer; cervical cancer; melanoma; leukaemia; TAT136.
XX
XX   OS   Homo sapiens.
XX
XX   FH   Key      Location/Qualifiers
XX   Peptide 1..28
XX   Protein 29..282
XX   /label= Signal_peptide
XX   /label= Mature_TAT136
XX   /note= "Tumour associated antigenic target polypeptide"
XX   Region 52..58
XX   /label= N-myristoylation_site
XX   Region 112..116
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XX   Region 126..132
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XX   Region 188..194
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XX   Region 205..209
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 FT Region 220..224
 FT /label= N-glycosylation_site
 FT Domain 258..282
 FT /label= Transmembrane_domain
 XX
 XX WO200216591-A2.
 XX
 XX 28-FEB-2002.
 XX
 XX 14-AUG-2001; 2001WO-US025464.
 XX
 XX 24-AUG-2000; 2000WO-US023328.
 XX
 XX 28-FEB-2001; 2001WO-US006520.
 PR
 PR 22-JUN-2001; 2001US-00888257.
 PR
 PR 22-JUN-2001; 2001WO-US020118.
 XX
 XX (GETH) GENENTECH INC.
 PA
 XX Gao W, Polakis P, Shou J, Smith V, Soriano R, Williams PM;
 PI Wu TD, Zhang Z;
 FI
 XX WPI; 2002-280928/32.
 DR
 DR N-PSDB; ABK11744.
 XX
 XX Novel isolated antibody which binds to tumor-associated antigenic target
 PT polypeptide useful for killing cancer cells expressing the polypeptide
 FT and for treating tumor comprising cells that expresses the polypeptide.
 FT
 XX
 XX Claim 2; Fig 8; 123pp; English.
 PS
 XX The invention describes an isolated antibody which binds to a tumour-
 CC associated antigenic target (TAT) polypeptide. The antibody is useful
 CC for: killing a cancer cell (such as a breast, colorectal, lung, ovarian,
 CC central nervous system, liver, bladder, pancreatic, cervical, melanoma or
 CC leukaemia cell) that expresses a polypeptide with at least 80% identity
 CC to the TAT polypeptide sequence; treating a tumour comprising cells that
 CC express a polypeptide with at least 80% identity to the TAT polypeptide
 CC sequence; determining the presence of a polypeptide having at least 80 %
 CC identity to the TAT polypeptide sequence in a sample suspected of
 CC containing the polypeptide; diagnosing the presence of a tumour in a
 CC mammal, and for antibody dependent enzyme mediated prodrug therapy
 CC (ADEPT). This is the amino acid sequence of the tumour associated
 CC antigenic target polypeptide (TAT) 136, described in the invention
 .XX
 SQ Sequence 282 AA;

Query Match 100.0%; Score 282; DB 5; Length 282;
 Best Local Similarity 100.0%; Pred. No. 2.8e-286; Mismatches 0; Indels 0; Gaps 0;
 Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy 121 QLTDAGTYKCYIITSKGGNANLEYKTFGAFSMPEVNVVDYNASSETLRCEAPRFPQPTVV 180
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 Db 181 WASQVDGAFSEVSNFSELSNENVTMKVSVLVYVNTYNTYSCMIENDIAKATGDIKV 240
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Qy 241 TSEIKRSHQLQLNSKASLCVSSFFAISWALLPLSPYLMK 282
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 Db 241 TSEIKRSHQLQLNSKASLCVSSFFAISWALLPLSPYLMK 282
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RESULT 10
 ABG95880
 ID ABG95880 standard; protein; 282 AA.
 XX
 XX AC ABG95880;
 XX
 XX 10-DEC-2002 (first entry)
 XX
 XX Human secreted/transmembrane protein PRO1291.
 DE
 .XX
 KW Human; secreted protein; transmembrane protein; antirheumatic;
 KW antiarthritic; osteopathic; sports-related joint problem;
 KW articular cartilage defect; osteoarthritis; rheumatoid arthritis.
 XX
 OS Homo sapiens.
 XX
 PN US2002119130-A1.
 XX
 XX 29-AUG-2002.
 PD
 XX
 PF 06-DEC-2001; 2001US-00006867.
 XX
 XX 29-OCT-1997; 97US-0063435P.
 PR 29-OCT-1997; 97US-0064215P.
 PR 22-APR-1998; 98US-0082797P.
 PR 29-APR-1998; 98US-0083495P.
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 PR 02-JUN-1998; 98US-0087759P.
 PR 04-JUN-1998; 98US-0088021P.
 PR 04-JUN-1998; 98US-0088029P.
 PR 10-JUN-1998; 98US-0088030P.
 PR 10-JUN-1998; 98US-0088734P.
 PR 10-JUN-1998; 98US-0088740P.
 PR 10-JUN-1998; 98US-0088811P.
 PR 10-JUN-1998; 98US-0088824P.
 PR 11-JUN-1998; 98US-0088825P.
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 PR 16-JUN-1998; 98US-0089105P.
 PR 17-JUN-1998; 98US-0089514P.
 PR 19-JUN-1998; 98US-0089653P.
 PR 22-JUN-1998; 98US-0089952P.
 PR 24-JUN-1998; 98US-0090246P.
 PR 25-JUN-1998; 98US-0090444P.
 PR 25-JUN-1998; 98US-0090688P.
 PR 26-JUN-1998; 98US-0090896P.
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 PR 17-AUG-1998; 98US-0096012P.
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 PR 01-SEP-1998; 98US-0097979P.
 PR 10-SEP-1998; 98US-0098749P.
 PR 10-SEP-1998; 98US-0099741P.
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 PR 10-SEP-1998; 98US-0099812P.
 PR 16-SEP-1998; 98US-0099815P.
 PR 16-SEP-1998; 98US-0100627P.
 PR 16-SEP-1998; 98US-0100662P.
 PR 16-SEP-1998; 98US-0100627P.
 PR 17-SEP-1998; 98US-0100684P.
 PR 17-SEP-1998; 98US-0100930P.
 PR 22-SEP-1998; 98US-0101279P.
 PR 23-SEP-1998; 98US-0101475P.
 PR 24-SEP-1998; 98US-0101738P.
 PR 24-SEP-1998; 98US-0101743P.
 PR 24-SEP-1998; 98US-0101916P.
 PR 30-SEP-1998; 98US-0102570P.
 PR 06-OCT-1998; 98US-0103449P.
 PR

FT Modified-site 205..209 /note= "Asn is N-glycosylated"
 FT Modified-site 216..220 /note= "Asn is N-glycosylated"
 FT Modified-site 220..224 /note= "Asn is N-glycosylated"
 FT Domain 258..281 /note= "Transmembrane domain"
 XX WO200216429-A2.
 XX 28-FEB-2002.
 XX 22-JUN-2001; 2001WO-US020118.
 XX 24-AUG-2000; 2000WO-US023328.
 PR 26-SEP-2000; 2000US-0235451P.
 PR 01-DEC-2000; 2000WO-US032678.
 PR 28-FEB-2001; 2001WO-US006520.
 PR 01-MAR-2001; 2001WO-US006666.
 XX (GETH) GENENTECH INC.
 XX Goddard A, Godowski PJ, Gurney AL, Hillan KJ, Polakis P, Smith V;
 PI Wood WI, Wu TD, Zhang Z;
 XX WPI; 2002-280917/32.
 DR N-PSDB; ABK11091.
 XX Novel isolated tumor-associated antigenic target polypeptides which are
 XX useful as targets for cancer therapy and diagnosis in mammals.
 XX Claim 12; Fig 8; 12pp; English.
 CC The invention relates to an isolated tumour-associated antigenic target
 CC polypeptide (TAT) (I), specifically TAT134-TAT138 polypeptides, and the
 CC polynucleotides (II) encoding them. (II) is useful for diagnosing the
 CC presence of a tumour in a mammal, where the level of expression of (II)
 CC is indicative on the presence of tumour in the mammal from which the test
 CC sample was obtained. Antibody to (I) is useful for killing a cancer cell,
 CC (e.g. breast cancer cell, a colorectal cancer cell, a lung cancer cell,
 CC an ovarian cancer cell, a central nervous system (CNS) cancer cell, a
 CC liver cancer cell, a bladder cancer cell, a pancreatic cancer cell, a
 CC melanoma cell or a leukaemia cell) that expresses (I). Oligonucleotides
 CC hybridising to (II) are useful as diagnostic probes, antisense
 CC oligonucleotide probes or for encoding fragments of full length TAT
 CC polypeptide. (II) is also useful in chromosome and gene mapping and in
 CC the generation of antisense RNA and DNA probes, for constructing
 CC hybridisation probes for mapping the gene encoding TAT and for genetic
 CC analysis of individuals with genetic disorders. (II) is also useful for
 CC generating either transgenic animals or knockout animals, and in gene
 CC therapy. The TAT polypeptides and nucleic acids may also be used for
 CC tissue typing and the TAT polypeptides are useful for screening compounds
 CC that mimic the TAT polypeptide (agonist) or prevent the effect of TAT
 CC polypeptide (antagonist). The antibody is useful for staging TAT
 CC polypeptide-expressing cancers, purifying or immunoprecipitating TAT
 CC polypeptide from cells, for detection and quantitation of TAT polypeptide
 CC in vitro, e.g., in an enzyme linked immunosorbent assay (ELISA) or
 CC western blot. The antibodies are also useful for treating a TAT-
 CC expressing cancer or alleviating one or more symptoms of cancer in a
 CC mammal. The present sequence represents the amino acid sequence of TAT136
 XX
 SQ Sequence 282 AA;
 Query Match 100.0%; Score 282; DB 5; Length 282;
 Best Local Similarity 100.0%; Pred. No. 2.8e-286;
 Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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 Db 1 MASLGQLFWISIIIIILAGALALIGFGISGRHSITVTTVASAGNIGDGLSCTFEP 60
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 QY 241 TESEIKRSHLQLLNKSKASLCVSPFAISWALLPLSPYMLK 282
 Db 241 TESEIKRSHLQLLNKSKASLCVSPFAISWALLPLSPYMLK 282
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 ID ABP30901 standard; protein; 282 AA.
 XX ABP30901;
 XX 02-JUL-2002 (first entry)
 DT OSE protein #2.
 DE Human; immunostimulant; cytostatic; cancer; ovarian carcinoma.
 XX Homo sapiens.
 OS Homo sapiens.
 PN WO200206317-A2.
 XX 24-JAN-2002.
 XX 17-JUL-2001; 2001WO-US022635.
 XX 17-JUL-2000; 2000US-00617747.
 PR 10-AUG-2000; 2000US-00636801.
 PR 20-SEP-2000; 2000US-00667857.
 PR 04-APR-2001; 2001US-00827271.
 PR 18-JUN-2001; 2001US-00884441.
 XX (CORI-) CORIXA CORP.
 XX Mitcham JL, King GE, Algate PA, Fling SP, Retter MW, Fanger GR;
 PI Reed SG, Vedvick TS, Carter D, Hill P, Albone E;
 XX WPI; 2002-164781/21.
 DR N-PSDB; ABN72971.
 XX Polypeptides comprising an immunogenic portion of an ovarian carcinoma
 XX protein or its variants, useful for stimulating an immune response in a
 XX patient and treating ovarian cancer.
 XX Claim 34; Page 321-322; 408pp; English.
 CC This invention relates to polypeptides comprising an immunogenic portion
 CC of an ovarian carcinoma protein which acts as an immunostimulant and is
 CC cytostatic. The polypeptides, polynucleotides, antibodies, fusion
 CC proteins, T cell populations and antigen presenting cells that express
 CC the polypeptides are useful for stimulating an immune response in a
 CC patient and treating ovarian cancer. This sequence represents protein
 CC related to the invention
 XX
 SQ Sequence 282 AA;
 Query Match 100.0%; Score 282; DB 5; Length 282;
 Best Local Similarity 100.0%; Pred. No. 2.8e-286;
 Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 MASLGQLFWISIIIIILAGALALIGFGISGRHSITVTTVASAGNIGDGLSCTFEP 60
 Db 1 MASLGQLFWISIIIIILAGALALIGFGISGRHSITVTTVASAGNIGDGLSCTFEP 60

(AMGE-) AMGEN INC.
 Fox M, Sullivan JK, Fang M;
 WPI; 2002-171639/22.
 N-PSDB; AAD29253.

Novel B7-like polypeptides, polynucleotides and their modulators useful for prevention and treatment of reproductive, immune and proliferative disorders, e.g. cancer, arteriosclerosis.

Claim 13; Fig 1A-1B; 133pp; English.

The present invention relates to an isolated B7-like (B7-L) polypeptide and its polynucleotide. B7-1 and its modulators are useful for treating reproductive disorders (e.g. infertility, miscarriage, preterm labour and delivery and endometriosis) and proliferative disorders. Antibodies, soluble proteins comprising extracellular domains and other regulators of B7-L are useful for enhancing the immune response to tumours. B7-1 plays a role in growth and maintenance of cancer cells based on the observation of seminal vesicle hyperplasia in transgenic mice overexpressing B7-1. Modulators of B7-1 are useful for the treatment of cancer e.g. seminal vesicle, lung, brain, breast, ovarian, testicular cancer and cancers of haematopoietic system. B7-1 and their modulators are useful to treat autoimmune diseases such as systemic lupus erythematosus, rheumatoid arthritis, immune thrombocytopenic purpura and psoriasis, chronic inflammatory disease such as inflammatory bowel disease (Crohn's disease and ulcerative colitis), Grave's disease, Hashimoto's thyroiditis and diabetes mellitus. They are also useful as immunosuppressive agents for bone marrow and organ transplantation or to prolong graft survival. Modulators of B7-L are also useful for diagnosis and treatment of diseases involving abnormal cell proliferation, arteriosclerosis and vascular restenosis. Soluble B7-L serves as vaccine adjuvants.

Antagonists of B7-L are useful for alleviation of toxic shock syndrome or allo sensitisation due to blood transfusions, and for treatment of multiple sclerosis, allergy, asthma and hypersensitivity reactions, nephropathies (e.g. glomerulonephritis), skin disorders (pemphigus and pemphigoid), endocrinopathies, various pneumopathies, vasculopathies, coeliac disease, anaemia, thrombocytopaenia, Gullain-Barre syndrome and myasthenia gravis, and lymphoproliferative disorders such as multiple myeloma. B7-L gene is useful in gene therapy and to map the locations of B7-L gene and related genes on chromosomes, as hybridisation probes in diagnostic assays, for isolating corresponding chromosomal B7-L genes, and to identify heritable tissue-degenerating diseases. The present sequence is human B7-L protein

Sequence 282 AA;

Query Match 100.0%; Score 282; DB 5; Length 282;
 Best Local Similarity 100.0%; Pred. No. 2.8e-286;
 Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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 DB 1 MASLGQLFWSIIISIIILAGAIALLIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP 60

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QY 241 TSEIKRSHQLQNLNSKASLCVSSFFAISWALLPLSPYMLK 282
 DB 241 TSEIKRSHQLQNLNSKASLCVSSFFAISWALLPLSPYMLK 282

RESULT 15

ABB09879
 ID ABB09879 standard; protein; 282 AA.
 XX
 AC ABB09879;
 XX
 DT 30-JUL-2002 (first entry)
 XX
 DE Amino acid sequence of the OREO gene (gene B).
 XX
 KW Human; gene A; ovarian tumour; gene B; OREO; ovarian cancer.
 XX
 OS Homo sapiens.
 XX
 FH Key Location/Qualifiers
 FT Domain 12..31
 FT /note= "predicted transmembrane domain"
 FT Domain 46..145
 FT /note= "predicted Ig domain"
 FT Modified-site 112
 FT /note= "N-glycosylation site"
 FT Modified-site 160
 FT /note= "N-glycosylation site"
 FT Modified-site 190
 FT /note= "N-glycosylation site"
 FT Modified-site 196
 FT /note= "N-glycosylation site"
 FT Modified-site 205
 FT /note= "N-glycosylation site"
 FT Modified-site 216
 FT /note= "N-glycosylation site"
 FT Modified-site 220
 FT /note= "N-glycosylation site"
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 WO2001194641-A2.
 PN
 PD 13-DEC-2001.
 XX
 PD 11-JUN-2001; 2001WO-US018700.
 XX
 PR 09-JUN-2000; 2000US-0210451P.
 XX
 PA (IDEC-) IDEC PHARM CORP.
 XX
 PI Ople E, McLachlan K, Heard C;
 XX
 DR WPI; 2002-404365/43.
 DR N-PSDB; ABL56582.
 XX
 PT New polynucleotide and corresponding antigens from human ovarian cancer cells, useful for treatment and diagnosis of ovarian cancer.
 PT
 XX
 PS Claim 12; Fig 7b; 71pp; English.
 XX
 CC The present sequence represents a protein designated OREO. The OREO (Ople designated gene B. This gene was identified by representational difference analysis (RDA) screening, and is selectively expressed by certain human ovarian tumours. The specification also describes gene A, identified by the same method. Gene A and B polynucleotides are useful for detecting ovarian cancer. Their polypeptides are useful for treating ovarian cancer
 CC
 CC Sequence 282 AA;

Query Match 100.0%; Score 282; DB 5; Length 282;
 Best Local Similarity 100.0%; Pred. No. 2.8e-286;
 Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MASLGQLFWSIIISIIILAGAIALLIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP 60
 DB 1 MASLGQLFWSIIISIIILAGAIALLIGFGISGRHSITVTTVASAGNIGEDGILSCTFEP 60

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Db	61	DIKLSDIVIOWLKEGVGLVHEFKEGKDELSEQDEMRGRTAVFADQVIVGNASRLKNV	120
Qy	121	QLTDAGTYKCYIITSKGKGNANLEYKTGAFSPMEVNVVDYNASSETLRCEAPRWFPOPTVV	180
Db	121	QLTDAGTYKCYIITSKGKGNANLEYKTGAFSPMEVNVVDYNASSETLRCEAPRWFPOPTVV	180
Qy	181	WASQVDOGANFSEVSNTSFELNSENVMTMKVSVLYNVVTINNTYSCMIENDIAKATGDIKV	240
Db	181	WASQVDOGANFSEVSNTSFELNSENVMTMKVSVLYNVVTINNTYSCMIENDIAKATGDIKV	240
Qy	241	TESEIKRRSHLOLLNSKASLCVSSFFFAISWALLPLSPYMLK	282
Db	241	TESEIKRRSHLOLLNSKASLCVSSFFFAISWALLPLSPYMLK	282

Search completed: April 19, 2005, 07:31:43
Job time : 92 secs

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GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: April 19, 2005, 07:17:40 ; Search time 83 Seconds
(without alignments)
1739.835 Million cell updates/sec

Title: US-10-773-715-6
Perfect score: 282
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Scoring table: OLIGO
Gapop 60.0 , Gapext 60.0

Searched: 1612378 seqs, 512079187 residues

Word size : 0

Total number of hits satisfying chosen parameters: 1612378

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Listing first 45 summaries

Database : UniProt_03.*

- 1: uniprot_sprot.*
- 2: uniprot_trembl.*

*Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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1	282	100.0	282	2 Q7Z7D3	Q7Z7D3 homo sapien
2	228	80.9	282	2 Q9H6B2	Q9H6B2 homo sapien
3	187	66.3	187	2 Q6P097	Q6P097 homo sapien
4	61	21.6	283	2 Q8K091	Q8K091 mus musculu
5	61	21.6	283	2 Q7TPH5	Q7TPH5 mus musculu
6	61	21.6	283	2 Q7TSP5	Q7TSP5 mus musculu
7	8	2.8	84	2 Q9B8D0	Q9B8D0 candida alb
8	8	2.8	106	2 Q8HDXD1	Q8HDXD1 macaca fasc
9	8	2.8	106	2 Q897X0	Q897X0 clostridium
10	8	2.8	118	2 Q6V1Y6	Q6V1Y6 pagrus majo
11	8	2.8	129	2 Q862X1	Q862X1 bos taurus
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13	8	2.8	145	2 Q862L6	Q862L6 bos taurus
14	8	2.8	153	2 Q68F82	Q68F82 xenopus tro
15	8	2.8	164	2 Q8C2K0	Q8C2K0 mus musculu
16	8	2.8	165	1 RL12_BOVIN	P61284 bos taurus
17	8	2.8	165	1 RL12_HUMAN	P30050 homo sapien
18	8	2.8	165	1 RL12_MOUSE	P35979 mus musculu
19	8	2.8	165	1 RL12_NEUCR	Q9C285 neurospora
20	8	2.8	165	1 RL12_RAT	P23358 rattus norv
21	8	2.8	165	2 Q608B6	Q608B6 homo sapien
22	8	2.8	165	2 Q6QM27	Q6QM27 chinchilla
23	8	2.8	165	2 Q66GW1	Q66GW1 xenopus lae
24	8	2.8	165	2 Q6DRB6	Q6DRB6 brachydanio
25	8	2.8	165	2 Q7ZUG1	Q7ZUG1 brachydanio
26	8	2.8	165	2 Q8AVW0	Q8AVW0 xenopus lae
27	8	2.8	167	2 Q90YV6	Q90YV6 italearius p
28	8	2.8	172	2 Q7US11	Q7US11 rhodopirell
29	8	2.8	174	2 Q7Q0Y7	Q7Q0Y7 anopheles g
30	8	2.8	191	2 Q6QAS5	Q6QAS5 sus scrofa
31	8	2.8	218	2 Q6DI58	Q6DI58 mus musculu

32	8	2.8	261	2 Q82UT7	Q82UT7 nitrosomona
33	8	2.8	279	2 Q7U3C8	Q7U3C8 synechococc
34	8	2.8	279	2 Q7V3T7	Q7V3T7 prochloroco
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36	8	2.8	299	1 HTPX_STRMU	Q93d93 streptococc
37	8	2.8	314	2 Q92TF9	Q92TF9 rhizobium m
38	8	2.8	370	1 YGP9_YEAST	P53110 saccharomyc
39	8	2.8	430	2 Q6ADS0	Q6ADS0 leifsonia x
40	8	2.8	564	2 Q6CW22	Q6CW22 kluyveromyc
41	8	2.8	569	2 Q7U928	Q7U928 synechococc
42	8	2.8	575	1 HEMA_P11HW	P16071 human parai
43	8	2.8	575	2 Q573I0	O57310 human parai
44	8	2.8	575	2 Q06992	Q06992 human parai
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ALIGNMENTS

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ID Q7Z7D3 PRELIMINARY; PRT; 282 AA.
AC Q7Z7D3;
DT 01-OCT-2003 (TrEMBLrel. 25, Created)
DT 01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
DT 25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
DE Immune costimulatory protein B7-H4 (T cell costimulatory molecule B7x) (B7h.5).
DE NCBI_TaxID=9606;
GN Name=B7-H4; ORFNames=UNQ659;
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RA Sica G.L., Choi I.-H., Zhu G., Tamada K., Wang S.-D., Tamura H., Chapoval A.I., Flies D.B., Bajorath J., Chen L.;
RL Submitted (APR-2003) to the EMBL/GenBank/DBJ databases.
RN [2]
RP SEQUENCE FROM N.A.
RX MEDLINE=22833980; PubMed=12920180; DOI=10.1073/pnas.1434299100;
RA Zang X., Loke P., Kim J., Murphy K., Waitz R., Allison J.P.;
RT "B7x: a widely expressed B7 family member that inhibits T cell activation.";
RL Proc. Natl. Acad. Sci. U.S.A. 100:10388-10392(2003).
RN [3]
RP SEQUENCE FROM N.A.
RX MEDLINE=22887296; PubMed=12975309; DOI=10.1101/gr.1293003;
RA Clark H.F., Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J., Chen J., Chow B., Chui C., Crowley C., Currell B., Deuel B., Dowd P., Eaton D., Foster J., Grimaldi C., Gu Q., Hass P.E., Heldens S., Huang A., Kim H.S., Klimowski L., Jin Y., Johnson S., Lee J., Lewis L., Liao D., Mark M., Robbie E., Sanchez C., Schoenfeld J., Seshagiri S., Simmons L., Singh J., Smith V., Stinson J., Vegts A., Vanden R., Watanabe C., Wiedand D., Woods K., Xie M.H., Yaneura D., Yi S., Yu G., Yuan J., Zhang M., Zhang Z., Goddard A., Wood W.I., Godowski P.;
RT "The secreted protein discovery initiative (SPDI), a large-scale effort to identify novel human secreted and transmembrane proteins: a bioinformatics assessment.";
RL Genome Res. 13:2265-2270(2003).
RN [4]
RP SEQUENCE FROM N.A.
RX TISSUE=Brain;
MEDLINE=22378257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Straubeberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D., Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K., Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F., Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L., Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E., Brownstein M.J., Udén T.B., Toshiyuki S., Carninci P., Prange C., Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

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RA Boesak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahney J., Helton E., Kettaman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smallus D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Marra M.A.;
RT "Generation and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences.";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
RN [5]
RP SEQUENCE FROM N.A.
RC TISSUE=Brain;
RA Director MGC Project;
RL Submitted (JUN-2004) to the EMBL/GenBank/DBJ databases.
DR EMBL: AY280972; AAB37283.1; -
DR EMBL: AY346100; AA024206.1; -
DR EMBL: AY358352; AAQ88718.1; -
DR EMBL: BC074729; AAH74729.1; -
DR HSP; Q63345; 1PKO.
DR InterPro; IPR007110; Ig-like.
DR Pfam; PF00047; Ig; 1.
DR PROSITE; PS50835; IG LIKE; 2.
SQ SEQUENCE 282 AA; 30878 MW; 1C9C565A9242E78C CRC64;

Query Match 100.0%; Score 282; DB 2; Length 282;
Best Local Similarity 100.0%; Pred. No. 5.6e-286;
Matches 282; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MASIGQILFWSIIIIIIILAGAIALLIIGFISGRHSITVTVASAGNIGDGILSCTFEP 60
Db 1 MASIGQILFWSIIIIIIILAGAIALLIIGFISGRHSITVTVASAGNIGDGILSCTFEP 60

Qy 61 DIKLSDIVIQWLKEGVLGVHFEKGDSEQDEMFRGRTAVFADQVIVGNASLRKNV 120
Db 61 DIKLSDIVIQWLKEGVLGVHFEKGDSEQDEMFRGRTAVFADQVIVGNASLRKNV 120

Qy 121 QLTDAQYKCYIITSKGKNANLEYKTCGAFSMPEVNVYDYNASSETLRCCEAPRPF 180
Db 121 QLTDAQYKCYIITSKGKNANLEYKTCGAFSMPEVNVYDYNASSETLRCCEAPRPF 180

Qy 181 WASQVQDQANFSEVNTSFELNSNVTKVSVLYNVTINNTYSCMIENDIAKATGDIKV 240
Db 181 WASQVQDQANFSEVNTSFELNSNVTKVSVLYNVTINNTYSCMIENDIAKATGDIKV 240

Qy 241 TESIIRKSHLQLNSKASLCVSSFFFAISWALLPLSPYLMK 282
Db 241 TESIIRKSHLQLNSKASLCVSSFFFAISWALLPLSPYLMK 282

RESULT 2
Q9H6B2
ID Q9H6B2 PRELIMINARY; PRT; 282 AA.
AC Q9H6B2
DT 01-MAR-2001 (TrEMBLrel. 16, Created)
DT 01-MAR-2001 (TrEMBLrel. 16; Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Hypothetical protein FLJ22418.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Prostate;
RA Strausberg R.;
RL Submitted (JAN-2004) to the EMBL/GenBank/DBJ databases.
RA Tanaka T., Nakamura Y., Isogai T., Sugano S.;
RL Submitted (AUG-2000) to the EMBL/GenBank/DBJ databases.
DR EMBL: AK026071; BAB15349.1; -
DR HSP; Q63345; 1PKO.
DR InterPro; IPR003599; Ig.

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DR InterPro; IPR007110; Ig-like.
DR SMART; SM00409; IG; 1.
DR PROSITE; PS50835; IG LIKE; 2.
SQ SEQUENCE 282 AA; 30893 MW; 6F9066999A1E9DB4 CRC64;

Query Match 80.9%; Score 228; DB 2; Length 282;
Best Local Similarity 100.0%; Pred. No. 1.7e-229;
Matches 228; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 55 SCTFEPDIKLSDIVIQWLKEGVLGVHFEKGDSEQDEMFRGRTAVFADQVIVGNAS 114
Db 55 SCTFEPDIKLSDIVIQWLKEGVLGVHFEKGDSEQDEMFRGRTAVFADQVIVGNAS 114

Qy 115 LRLKNVLTDAQYKCYIITSKGKNANLEYKTCGAFSMPEVNVYDYNASSETLRCCEAPRWF 174
Db 115 LRLKNVLTDAQYKCYIITSKGKNANLEYKTCGAFSMPEVNVYDYNASSETLRCCEAPRWF 174

Qy 175 PQPTVWASQVQDQANFSEVNTSFELNSNVTKVSVLYNVTINNTYSCMIENDIAKA 234
Db 175 PQPTVWASQVQDQANFSEVNTSFELNSNVTKVSVLYNVTINNTYSCMIENDIAKA 234

Qy 235 TGDIKVTESEIKRSHLQLNSKASLCVSSFFFAISWALLPLSPYLMK 282
Db 235 TGDIKVTESEIKRSHLQLNSKASLCVSSFFFAISWALLPLSPYLMK 282

RESULT 3
Q6P097
ID Q6P097 PRELIMINARY; PRT; 187 AA.
AC Q6P097
DT 05-JUL-2004 (TrEMBLrel. 27, Created)
DT 05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
DT 05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
DE B7-H4 protein.
GN Name=B7-H4;
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Prostate;
RX MEDLINE=22398257; Pubmed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Dege J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Heish F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullaby S.J.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahney J., Helton E., Kettaman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smallus D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Marra M.A.;
RT "Generation and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences.";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
RN [2]
RP SEQUENCE FROM N.A.
RC TISSUE=Prostate;
RA Strausberg R.;
RL Submitted (JAN-2004) to the EMBL/GenBank/DBJ databases.
DR EMBL: BC065717; AAB65717.1; -
DR InterPro; IPR007110; Ig-like.
DR PROSITE; PS50835; IG LIKE; 1.
SQ SEQUENCE 187 AA; 20743 MW; 9AED6155AEBD4E63 CRC64;

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Query Match 66.3%; Score 187; DB 2; Length 187;
 Best Local Similarity 100.0%; Pred. No. 9.4e-187;
 Matches 187; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 96 MFRGRTAVFADQVIVGNASRLKXVQLTDAGTYKCYIITSKGNANLEYKTAFAFSPFV 155
 Db 1 MFRGRTAVFADQVIVGNASRLKXVQLTDAGTYKCYIITSKGNANLEYKTAFAFSPFV 60

Qy 156 NVDYNASSETLRCEAPRWFPQPTVWASQVDOGANFSEVNTSPFELNSVNTSKVSVLY 215
 Db 61 NVDYNASSETLRCEAPRWFPQPTVWASQVDOGANFSEVNTSPFELNSVNTSKVSVLY 120

Qy 216 NVTINNTYSCMIENDIAKATGDIKVTSEIKRSHLQLLNKASLCVSSPFAISWALLPL 275
 Db 121 NVTINNTYSCMIENDIAKATGDIKVTSEIKRSHLQLLNKASLCVSSPFAISWALLPL 180

Qy 276 SPYLMK 282
 Db 181 SPYLMK 187

RESULT 4
 Q8K091 PRELIMINARY; PRT; 283 AA.
 AC Q8K091
 DT 01-OCT-2002 (TrEMBLrel. 22, Created)
 DT 01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
 DT 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
 DE Immune costimulatory protein B7-H4.
 GN Name=BC032925;
 OS Mus musculus (Mouse).
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
 OX NCBI_TaxID=10090;
 RN [1]
 RP SEQUENCE FROM N.A.
 RC TISSUE=Uterus;
 RX MEDLINE=22389257; PubMed=12477932; DOI=10.1073/pnas.242603899;
 RA Strauberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
 RA Klausner R.D., Collins F.S., Wagner L., Shennan C.M., Schuler G.D.,
 RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
 RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Haieh F.,
 RA Datchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
 RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.B., Scheetz T.E.,
 RA Brownstein M.J., Udén T.B., Toshiyuki S., Carninci P., Prange C.,
 RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
 RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
 RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
 RA Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
 RA Fahy J., Helton E., Kettelman M., Madan A., Rodrigues S., Sanchez A.,
 RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
 RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
 RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
 RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
 RA Jones S.J., Marra M.A.;
 RA "Generation and initial analysis of more than 15,000 full-length human
 RT and mouse cDNA sequences.";
 RL Proc. Natl. Acad. Sci. U.S.A. 99:16999-16903 (2002).
 RN [2]
 RP SEQUENCE FROM N.A.
 RC TISSUE=Uterus;
 RA Strauberg R.;
 RL Submitted (JUN-2002) to the EMBL/GenBank/DBJ databases.
 DR EMBL; BC032925; AAH32925.1; -;
 DR HSP; Q63345; IPKO
 DR MGD; MGI:3039619; BC032925.
 DR InterPro; IPR003599; IG.
 DR InterPro; IPR007110; IG-like.
 DR SMART; SM00409; IG; 1.
 DR PROSITE; PS50835; IG LIKE; 2.
 SQ SEQUENCE 283 AA; 30801 MW; 755817417323453B CRC64;

Query Match 21.6%; Score 61; DB 2; Length 283;

Best Local Similarity 100.0%; Pred. No. 9e-55;
 Matches 61; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 181 WASQVDOGANFSEVNTSPFELNSVNTSKVSVLYNTYSCMIENDIAKATGDIKV 240
 Db 181 WASQVDOGANFSEVNTSPFELNSVNTSKVSVLYNTYSCMIENDIAKATGDIKV 240

Qy 241 T 241
 Db 241 T 241

RESULT 5
 Q7TPH5 PRELIMINARY; PRT; 283 AA.
 AC Q7TPH5
 DT 01-OCT-2003 (TrEMBLrel. 25, Created)
 DT 01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
 DT 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
 DE B7S1.
 OS Mus musculus (Mouse).
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
 OX NCBI_TaxID=10090;
 RN [1]
 RP SEQUENCE FROM N.A.
 RX MEDLINE=22703430; PubMed=12818166; DOI=10.1016/S1074-7613(03)00147-X;
 RA Prasad D.V., Richards S., Mai X.M., Dong C.;
 RA "B7S1, a novel B7 family member that negatively regulates T cell
 RT activation.";
 RL Immunity 18:863-873 (2003).
 DR EMBL; AY322147; AAP8965.1; -;
 DR HSP; Q63345; IPKO.
 DR InterPro; IPR003599; IG.
 DR SMART; SM00409; IG; 1. IG-like.
 DR PROSITE; PS50835; IG LIKE; 2.
 SQ SEQUENCE 283 AA; 30847 MW; A97F17461857850B CRC64;

Query Match 21.6%; Score 61; DB 2; Length 283;
 Best Local Similarity 100.0%; Pred. No. 9e-55;
 Matches 61; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 181 WASQVDOGANFSEVNTSPFELNSVNTSKVSVLYNTYSCMIENDIAKATGDIKV 240
 Db 181 WASQVDOGANFSEVNTSPFELNSVNTSKVSVLYNTYSCMIENDIAKATGDIKV 240

Qy 241 T 241
 Db 241 T 241

RESULT 6
 Q7TSP5 PRELIMINARY; PRT; 283 AA.
 AC Q7TSP5
 DT 01-OCT-2003 (TrEMBLrel. 25, Created)
 DT 01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
 DT 05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
 DE Immune costimulatory protein B7-H4 (T cell costimulatory molecule
 DE B7x).
 OS Mus musculus (Mouse).
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
 OX NCBI_TaxID=10090;
 RN [1]
 RP SEQUENCE FROM N.A.
 RC STRAIN=BALB/c;
 RA Sica G.L., Choi I.-H., Zhu G., Tamada K., Wang S.-D., Tamura H.,
 RA Chapoval A.I., Flies D.B., Bajorath J., Chen L.;
 RA Submitted (APR-2003) to the EMBL/GenBank/DBJ databases.
 RN [2]
 RP SEQUENCE FROM N.A.

QY	15	IIIIILAGA	22
Db	54	IIIIILAGA	61
RESULT 8			
Q8HXD1		PRELIMINARY;	PRT; 106 AA.
ID	Q8HXD1		
AC	Q8HXD1		
DT	01-MAR-2003	(TREMELrel. 23, Created)	
DT	01-MAR-2003	(TREMELrel. 23, Last sequence update)	
DT	01-MAR-2003	(TREMELrel. 23, Last annotation update)	
DE	Hypothetical protein.		
OS	Macaca fascicularis	(Crab eating macaque) (Cynomolgus monkey).	
OC	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;		
OC	Mammalia; Eutheria; Primates; Catarrhini; Cercopithecoidea;		
OC	Cercopithecinae; Macaca.		
OX	NCBI_TaxID=9541;		
OX	[1]		
RP	SEQUENCE FROM N.A.		
RC	TISSUE=Frontal lobe left;		
RC	MEDLINE=21458551; PubMed=11574149; DOI=10.1016/S0378-1119(01)00665-5;		
RC	Osada N., Hida M., Kusuda J., Tanuma R., Iseki K., Hirata M., Suto Y.,		
RA	Hirai M., Terao K., Suzuki Y., Sugano S., Hashimoto K.;		
RA	"Assignment of 118 novel cDNAs of cynomolgus monkey brain to human		
RT	chromosomes.";		
RT	Gene 275:31-37(2001).		
RL	[2]		
RP	SEQUENCE FROM N.A.		
RC	TISSUE=Frontal lobe left;		
RC	Hashimoto K., Osada N., Hida M., Kusuda J., Sugano S.;		
RL	Submitted (OCT-2002). to the EMBL/GenBank/DBJ databases.		
RL	EMBL; AB093659; BAC21633.1; --		
KW	Hypothetical protein.		
QY	SEQUENCE 106 AA; 11676 MW; 6F6420F390EA53C3 CRC64;		
		2.8%; Score 8; DB 2; Length 106;	
	Query Match		
	Best Local Similarity 100.0%; Pred. No. 10;		
	Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps		
QY	38	TVTTTASA	45
Db	28	TVTTTASA	35
RESULT 9			
Q897X0		PRELIMINARY;	PRT; 106 AA.
ID	Q897X0		
AC	Q897X0		
DT	01-JUN-2003	(TREMELrel. 24, Created)	
DT	01-JUN-2003	(TREMELrel. 24, Last sequence update)	
DT	01-MAR-2004	(TREMELrel. 26, Last annotation update)	
DE	Putative transcriptional regulator (Padr family).		
GN	Name=padr; OrderedLocusNames=CTC00601;		
OS	Clostridium tetani.		
OC	Bacteria; Firmicutes; Clostridia; Clostridiales; Clostridiaceae;		
OC	Clostridium.		
OX	NCBI_TaxID=1513;		
OX	[1]		
RP	SEQUENCE FROM N.A.		
RC	STRAIN=Massachusetts / B88;		
RC	MEDLINE=22457253; PubMed=12552429; DOI=10.1073/pnas.0335853100;		
RA	Brueggemann H., Baumer S., Fricke W.F., Wierze A., Liesegang H.,		
RA	Decker I., Herzberg C., Martinez-Arias R., Merkl R., Henne A.,		
RA	Gottschalk G.;		
RT	"The genome sequence of Clostridium tetani, the causative agent of		
RT	tetanus disease.";		
RL	Proc. Natl. Acad. Sci. U.S.A. 100:1316-1321(2003).		
DR	EMBL; AE015938; AA035216.1; --		
DR	InterPro; IPR005149; Padr.		
DR	InterPro; IPR009058; Wing_hlx_DNA_bnd.		
KW	Pfam; PF03551; Padr; 1.		
KW	Complete proteome.		

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SQ SEQUENCE 106 AA; 12356 MW; 2D2ECFC628204CDF CRC64;
Query Match 2.8%; Score 8; DB 2; Length 106;
Best Local Similarity 100.0%; Pred. No. 10;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 85 EKGDELSE 92
Db 81 EKGDELSE 88
|||||

RESULT 10
Q6YIY6 PRELIMINARY; PRT; 118 AA.
ID Q6YIY6
AC Q6YIY6
DT 05-JUL-2004 (TrEMBLrel. 27, Created)
DT 05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
DT 05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
DE Ribosomal protein L12 (Fragment).
OS Pegrus major (Red sea bream) (Chrysophrys major).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
OC Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes; Percoidae;
OC Sparidae; Pegrus.
OX NCBI_TaxID=143350;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Spleen;
RA Chen S.L., Xu M.Y.;
RL Submitted (DEC-2002) to the EMBL/GenBank/DBJ databases.
CC -i- SIMILARITY: Belongs to the ribosomal protein L12P family.
DR EMBL; AY190964; AAC92747.1; -.
DR GO; GO:0005840; C:ribosome; IEA.
DR GO; GO:0003735; F:structural constituent of ribosome; IEA.
DR GO; GO:0006412; P:protein biosynthesis; IEA.
DR InterPro; IPR000911; Ribosomal_L12.
DR Pfam; PF00298; Ribosomal_L12.
DR SMART; SM00649; RL11; 1.
DR Ribonucleoprotein; Ribosomal protein.
FT NON_TER 118
FT NON_TER 118
SQ SEQUENCE 118 AA; 12908 MW; 5C7A2F03D2DDA8FD CRC64;

Query Match 2.8%; Score 8; DB 2; Length 118;
Best Local Similarity 100.0%; Pred. No. 11;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 230 DIAKATGD 237
Db 15 DIAKATGD 22
|||||

RESULT 11
Q862X1 PRELIMINARY; PRT; 129 AA.
ID Q862X1
AC Q862X1
DT 01-JUN-2003 (TrEMBLrel. 24, Created)
DT 01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Similar to ribosomal protein L12 (Fragment).
OS Bos taurus (Bovine).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;
OC Bovinae; Bos.
OX NCBI_TaxID=9913;
RN [1]
RP SEQUENCE FROM N.A.
RC MEDLINE=22544902; PubMed=12658628; DOI=10.1002/mrd.10292;
RA Ishiwa H., Katsuna S., Kizaki K., Patel O.V., Nakano H.,
RA Takahashi T., Inai K., Hirasawa A., Shiojima S., Ikawa H., Suzuki Y.,
RA Tsujimoto G., Izaike Y., Todoroki J., Hashizume K.;
RT "Characterization of gene expression profiles in early bovine
pregnancy using a custom cDNA microarray.";
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RL Mol. Reprod. Dev. 65:9-18(2003).
CC -i- SIMILARITY: Belongs to the ribosomal protein L12P family.
DR EMBL; AB098784; BAC56320.1; -.
DR GO; GO:0005840; C:ribosome; IEA.
DR GO; GO:0003735; F:structural constituent of ribosome; IEA.
DR GO; GO:0006412; P:protein biosynthesis; IEA.
DR InterPro; IPR000911; Ribosomal_L12.
DR Pfam; PF00298; Ribosomal_L12.
DR Pfam; PF03946; Ribosomal_L12_N; 1.
DR SMART; SM00649; RL11; 1.
DR PROSITE; PS00359; RIBOSOMAL_L12; 1.
KW Ribonucleoprotein; Ribosomal protein.
FT NON_TER 129
FT NON_TER 129
SQ SEQUENCE 129 AA; 14127 MW; E9F90F265C3EC763 CRC64;

Query Match 2.8%; Score 8; DB 2; Length 129;
Best Local Similarity 100.0%; Pred. No. 12;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 230 DIAKATGD 237
Db 9 DIAKATGD 16
|||||

RESULT 12
O66334 PRELIMINARY; PRT; 141 AA.
ID O66334
AC O66334
DT 01-AUG-1998 (TrEMBLrel. 07, Created)
DT 01-AUG-1998 (TrEMBLrel. 07, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Alpha subunit of dinitrogenase reductase (Fe protein) (Fragment).
GN Name=nifH;
OS unidentified nitrogen-fixing bacteria.
OC Bacteria.
OX NCBI_TaxID=34107;
RN [1]
RP SEQUENCE FROM N.A.
RC MEDLINE=20011231; PubMed=10543805;
RA Okuma M., Noda S., Kudo T.;
RT "Phylogenetic diversity of nitrogen fixation genes in the symbiotic
microbial community in the gut of diverse termites."
RL Appl. Environ. Microbiol. 65:4926-4934(1999).
CC -i- SIMILARITY: Belongs to the nifH / bchL / chlL family.
DR EMBL; AB011904; BAA28439.1; -.
DR HSP; P00456; 1CP2.
DR GO; GO:0005524; F:ATP binding; IEA.
DR GO; GO:0016491; F:oxidoreductase activity; IEA.
DR GO; GO:0006118; P:electron transport; IEA.
DR InterPro; IPR000392; NitrogenaseII.
DR Pfam; PF00142; Fer4_NifH; 1.
DR PRINTS; PR00091; NITROGNASEII.
DR PROSITE; PS00746; NIFH_FRXC_1; 1.
KW 4Fe-4S; ATP-binding; Iron; Iron-sulfur; Metal-binding; Oxidoreductase.
FT NON_TER 141
FT NON_TER 141
SQ SEQUENCE 141 AA; 14987 MW; 15AAADCEB162617C CRC64;

Query Match 2.8%; Score 8; DB 2; Length 141;
Best Local Similarity 100.0%; Pred. No. 13;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 61 DIKLSDIV 68
Db 52 DIKLSDIV 59
|||||

RESULT 13
Q862L6 PRELIMINARY; PRT; 145 AA.
ID Q862L6
AC Q862L6;
DT 01-JUN-2003 (TrEMBLrel. 24, Created)
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DT 01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
 DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
 DE Similar to ribosomal protein L12 (Fragment).
 OS Bos taurus (Bovine).
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 OC Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;
 OC Bovinae; Bos.
 OX NCBI_TaxID=9913;
 RN [1]
 RP SEQUENCE FROM N.A.
 RX MEDLINE=22544902; PubMed=12658628; DOI=10.1002/mrd.10292;
 RA Ishiwata H., Katsuma S., Kizaki K., Patel O.V., Nakano H.,
 RA Takahashi T., Imai K., Hirasawa A., Shiojima S., Ikawa H., Suzuki Y.,
 RA Teujimoto G., Izaike Y., Todoroki J., Hashizume K.;
 RT "Characterization of gene expression profiles in early bovine
 RT pregnancy using a custom cDNA microarray.";
 RL Mol. Reprod. Dev. 65:9-18(2003).
 CC -1- SIMILARITY: Belongs to the ribosomal protein L11P family.
 DR EMBL; AB098966; BAC56456.1; -;
 DR HSP; P29395; 1MWS.
 DR GO; GO:0005840; C:ribosome; IEA.
 DR GO; GO:0003735; F:structural constituent of ribosome; IEA.
 DR GO; GO:0006412; P:protein biosynthesis; IEA.
 DR InterPro; IPR000911; Ribosomal_L11.
 DR Pfam; PF00298; Ribosomal_L11; 1.
 DR Pfam; PF03946; Ribosomal_L11_N; 1.
 DR SMART; SM00649; RL11; 1.
 DR PROSITE; PS00359; RIBOSOMAL_L11; 1.
 KW Ribonucleoprotein; Ribosomal protein.
 FT NON_TER 1
 FT NON_TER 145
 SQ SEQUENCE 145 AA; 15690 MW; 386D68BD728102D3 CRC64;

 Query Match 2.8%; Score 8; DB 2; Length 145;
 Best Local Similarity 100.0%; Pred. No. 14;
 Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

 QY 230 DIAKATGD 237
 DB |||||
 45 DIAKATGD 52

 RESULT 14
 Q68F82 PRELIMINARY; PRT; 153 AA.
 ID Q68F82
 AC Q68F82
 DT 25-OCT-2004 (TrEMBLrel. 28, Created)
 DT 25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
 DE Hypothetical protein.
 OS Xenopus tropicalis (Western clawed frog) (Silurana tropicalis).
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 OC Amphibia; Batrachia; Anura; Mesobatrachia; Pipidae;
 OC Xenopodinae; Xenopus.
 OX NCBI_TaxID=8364;
 RN [1]
 RP SEQUENCE FROM N.A.
 RC TISSUE=Embryo;
 RX PubMed=12477932; DOI=10.1073/pnas.242603899;
 RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
 RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
 RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
 RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Haieh F.,
 RA Diatchenko L., Narusina K., Farmer A.A., Rubin G.M., Hong L.,
 RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
 RA Brownstein M.J., Udén T.B., Toshiyuki S., Carninci P., Prange C.,
 RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullany S.J.,
 RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
 RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
 RA Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
 RA Fahey J., Helton E., Kettelman M., Madan A., Rodrigues S., Sanchez A.,
 RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
 RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
 RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
 RA Jones S.J., Marra M.A.;
 RT "Generation and initial analysis of more than 15,000 full-length human
 RT and mouse cDNA sequences.";
 RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
 RN [2]
 RP SEQUENCE FROM N.A.
 RC TISSUE=Embryo;
 RA Klein S., Gerhard D.S.;
 RL Submitted (Aug-2004) to the EMBL/GenBank/DBJ databases.
 CC -1- SIMILARITY: Belongs to the ribosomal protein L11P family.
 DR EMBL; BC079961; AAH79961.1; -;
 DR InterPro; IPR000911; Ribosomal_L11.
 DR Pfam; PF00298; Ribosomal_L11; 1.
 DR Pfam; PF03946; Ribosomal_L11_N; 1.
 DR SMART; SM00649; RL11; 1.
 DR PROSITE; PS00359; RIBOSOMAL_L11; 1.
 KW Hypothetical protein; Ribonucleoprotein; Ribosomal protein.
 SQ SEQUENCE 153 AA; 16408 MW; 51031F4215E29158 CRC64;

 Query Match 2.8%; Score 8; DB 2; Length 153;
 Best Local Similarity 100.0%; Pred. No. 14;
 Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

 QY 230 DIAKATGD 237
 DB |||||
 33 DIAKATGD 40

 RESULT 15
 Q8C2K0 PRELIMINARY; PRT; 164 AA.
 ID Q8C2K0
 AC Q8C2K0
 DT 01-MAR-2003 (TrEMBLrel. 23, Created)
 DT 01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
 DT 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
 DE Mus musculus 2 days neonate thymus thymic cells cDNA, RIKEN full-
 DE length enriched library, clone:B430018F03 product:ribosomal protein
 DE L12, full insert sequence.
 GN Name=Rpl12;
 OS Mus musculus (Mouse).
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Mus.
 OX NCBI_TaxID=10090;
 RN [1]
 RP SEQUENCE FROM N.A.
 RC STRAIN=NOD; TISSUE=Thymus;
 RX MEDLINE=99279253; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9;
 RA Carninci P., Hayashizaki Y.;
 RT "High-efficiency full-length cDNA cloning.";
 RL Meth. Enzymol. 303:19-44(1999).
 RN [2]
 RP SEQUENCE FROM N.A.
 RC STRAIN=NOD; TISSUE=Thymus;
 RX MEDLINE=21085660; PubMed=11217851; DOI=10.1038/35055500;
 RA RIKEN FANTOM Consortium;
 RT "Functional annotation of a full-length mouse cDNA collection.";
 RL Nature 409:685-690(2001).
 RN [3]
 RP SEQUENCE FROM N.A.
 RC STRAIN=NOD; TISSUE=Thymus;
 RA The FANTOM Consortium;
 RA the RIKEN Genome Exploration Research Group Phase I & II Team;
 RT "Analysis of the mouse transcriptome based on functional annotation of
 RT 60,770 full-length cDNAs.";
 RL Nature 420:563-573(2002).
 RN [4]
 RP SEQUENCE FROM N.A.
 RC STRAIN=NOD; TISSUE=Thymus;
 RX MEDLINE=20499374; PubMed=11042159; DOI=10.1101/gr.145100;
 RA Carninci P., Shibata Y., Hayatsu N., Sugahara Y., Shibata K., Itoh M.,
 RA Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.;

RT "Normalization and subtraction of cap-trapper-selected cDNAs to
 RL prepare full-length cDNA libraries for rapid discovery of new genes."
 RN Genome Res. 10:1617-1630(2000).

[5]

RN SEQUENCE FROM N.A.

RP STRAIN=NOD; TISSUE=Thymus;

RC MEDLINE=20530913; PubMed=11076861; DOI=10.1101/gr.152600;

RA Shibata K., Itoh M., Aizawa K., Nagaoka S., Sasaki N., Carninci P.,
 RA Konno H., Akiyama J., Nishi K., Kitsunai T., Tashiro H., Itoh M.,
 RA Sumi N., Ishii Y., Nakamura S., Hazama M., Nishine T., Harada A.,
 RA Yamamoto R., Matsumoto H., Sakaguchi S., Ikegami T., Kashiwagi K.,
 RA Fujiwaka S., Inoue K., Togawa Y., Izawa M., Ohara E., Watahiki M.,
 RA Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuura S., Kawai J.,
 RA Okazaki Y., Muramatsu M., Inoue Y., Kira A., Hayashizaki Y.,
 RT "RIKEN integrated sequence analysis (RISA) system-384-format
 RL sequencing pipeline with 384 multicapillary sequencer."
 RN Genome Res. 10:1757-1771(2000).

[6]

RN SEQUENCE FROM N.A.

RP STRAIN=NOD; TISSUE=Thymus;

RC Adachi J., Aizawa K., Akimura T., Arakawa T., Bono H., Carninci P.,
 RA Fukuda S., Furuno M., Hanagaki T., Hara A., Hashizume W.,
 RA Hayashida K., Hayatsu N., Hiramoto K., Hiraoka T., Hirozane T.,
 RA Hori F., Imotani K., Ishii Y., Itoh M., Kagawa I., Kasukawa T.,
 RA Katoh H., Kawai J., Kojima Y., Kondo S., Konno H., Kouda M., Koya S.,
 RA Kurihara C., Matsuyama T., Miyazaki A., Murata M., Nakamura M.,
 RA Nishi K., Nomura K., Numazaki R., Ohno M., Ohsato N., Okazaki Y.,
 RA Saito R., Saitoh H., Sakai C., Sakai K., Sakazume N., Sano H.,
 RA Sasaki D., Shibata K., Shinagawa A., Shiraki T., Sogabe Y., Tagami M.,
 RA Tagawa A., Takahashi F., Takaku-Akahira S., Takeda Y., Tanaka T.,
 RA Tomaru A., Toya T., Yasunishi A., Muramatsu M., Hayashizaki Y.,
 RL Submitted (APR-2002) to the EMBL/GenBank/DBJ databases.
 CC -1- SIMILARITY: Belongs to the ribosomal protein L11p family.
 DR EMBL; AK088464; BAC40369.1; --

DR HSSP; P29195; 1MWS.

DR MGD; MGJ:98002; Rpl12.

DR GO; GO:0005737; C:cytoplasm; IDA.

DR GO; GO:0005730; C:nucleolus; IDA.

DR GO; GO:0005515; F:protein binding; IPI.

DR InterPro; IPR000911; Ribosomal_L11.

DR Pfam; PF00298; Ribosomal_L11; 1.

DR Pfam; PF03946; Ribosomal_L11_N; 1.

DR SMART; SM00649; RL11; 1.

DR PROSITE; PS00359; RIBOSOMAL_L11; 1.

KW Ribonucleoprotein; Ribosomal protein.

SQ SEQUENCE 164 AA; 17806 MW; 66A010247E426D0C CRC64;

Query Match 2.8%; Score 8; DB 2; Length 164;

Best Local Similarity 100.0%; Pred.No. 15;

Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 230 DIAKATGD 237

Db 45 DIAKATGD 52

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Job time : 88 secs

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